

# Research trends in pharmacological modulation of tumor-associated macrophages

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### HIGHLIGHT

- The biofunctions and clinical implications of TAMs in cancer progression were summarized.
- The current pharmaceutical targets in TAMs regulation were introduced and discussed.



The biofunctions and clinical implications of TAMs in cancer progression as well as the current pharmaceutical targets in TAMs regulation were summarized. The latest advancements of agents that were effective in TAMs depletion or reprogramming were summarized and discussed. The promising future of exosomal signaling for TAMs-targeted therapy was discussed and expected.

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#### REVIEW



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Abbreviations: TAMs, tumor-associated macrophages; TME, tumor microenvironment; PAMP, pathogen-associated patterns; LPS, lipopolysaccharides; TNF-α, tumor necrosis factor α; GM-CSF, granulocyte-macrophage colony-stimulating factor; IL, interleukin; DAMP, damage-associated patterns; ARG1, arginase; DMBA, 12-dimethylbenz-(a)anthracene; sCD163, soluble CD163; irAEs, immune-related adverse events; VEGFA, vascular endothelial growth factor A; ADM, adrenomedullin; bFGF, basic fibroblast growth factor; PDGF, platelet-derived growth factor; MIF, macrophage-inhibitory factor; PAF, platelet-activating factor; monocyte chemoattractant protein-1 MCP-1, monocyte chemoattractant protein-1; MMPs, matrix metalloproteinases; PD-L1, program cell death ligand 1; CTLA-4, cytotoxic T-lymphocyte-associated protein 4; OS, overall survival; DFS, disease free survival; PSGL-1, P-selectin glycoprotein ligand-1; ICAM-1, intracellular cell adhesion molecule-1; ERK5, extracellular-regulated protein kinase 5; EMT, epithelial-mesenchymal transition; ECM, extracellular matrix; PMN, premetastatic niche; GPCRs, G-protein-coupled receptors; MDSCs, myeloid-derived suppressor cells; CTCL, cutaneous T-cell lymphoma; IFNs, interferons; RTKs, Tyrosine kinase receptors; CSF-1R, colony stimulation factor-1 receptor; MSP, macrophage-stimulating protein; MerTK, Mer tyrosine kinase; HIF, Hypoxia-inducible factors; BMDMs, bone marrow-derived macrophages; PPAR, peroxisome proliferator-activated receptor; PDK, pyruvate dehydrogenase kinase; IDO, indoleamine-2, 3-dioxygenase; RAGE, receptor for advanced glycation end-product; DAG, diacylglycerol; MG, methylglyoxal; AGEs, advanced glycation end products; ELISA, enzyme-linked immunosorbent assays; dt-GCTs, diffuse-type tenosynovial giant-cell tumors; exoASO, exosome carrying antisense oligonucleotide; TRL7/8, toll-like receptor 7 and 8; SIRP $\alpha$ , signal regulatory protein- $\alpha$ ; TREM2, triggering receptor expressed on myeloid cells 2; ER $\alpha$ , estrogen receptor  $\alpha$ ; ERK, extracellular signal-regulated kinase; PKC, protein kinase C; PI3K, phosphoinositide-3 kinase; AKT, the serine/threonine kinase; MAPK, mitogen-activated protein kinase; mTOR, mammalian/mechanistic target of rapamycin; NF-xB, nuclear factor xB; PDGF-BB, platelet-derived growth factor subunit B homodimer; Ezh2, enhancer of zeste homolog 2; CSCs, cancer stem cells; ECs, endothelial cells; STAT3, signal transducer and activator of transcription 3; CSN5, COP9 signalosome subunit 5; FOXP3, forkhead box P3; KLF5, kruppel-like factor 5; TGF-β, transforming growth factor  $\beta$ ; GSK-3 $\beta$ , Glycogen synthase kinase-3 $\beta$ ; ANXA2, annexin A2; PITPNM3, phosphatidylinositol transfer protein 3; FOXO1, Forkhead box O1; CEBPB, CCAAT/enhancer binding protein beta; TRAIL, tumor necrosis factor-related apoptosis-inducing ligand; ABCB1, ATP binding cassette subfamily B member 1; HuR, human antigen R; JAK2, Janus protein tyrosine kinases; Pyk2, proline-rich tyrosine kinase 2; ELMO1, engulfment and cell motility 1; SAPK, stress-activated protein kinase; JNK, Jun N-terminal kinase; HDAC1, histone deacetylase 1; TACE, tumor necrosis factor alpha-converting enzyme; SOX4, SRY-box transcription factor 4; S100A8/9, S100 calcium binding protein A8/9; TAK1, transforming growth factor-beta-activated kinase 1; EGFR, epidermal growth factor receptor; FAK, focal adhesion kinase; RhoA, ras homolog family member A; PGE2, Prostaglandin E2; ARHGAP10, Rho GTPase Activating Protein 10; MALAT1, metastasis-associated lung adenocarcinoma transcript 1; TOPK, T-LAK Cell-originated Protein Kinase; JNK, c-Jun N-terminal kinase; MST1R, macrophage-stimulating 1 receptor; HK2, hexokinase 2; GPR132, G protein-coupled receptor 132; REDD1, DNA damage response-1; HMGB1, high-mobility group box 1; RIPK3, receptor-interacting protein kinase-3; FABP, fatty acid binding protein; MCAD, medium-chain acyl-coA dehydrogenase; GS, glutamine synthetase; ARG, arginase; FPN, ferroportin; LCN, lipocalin; BRG1, brahma-related gene 1; LIPA, lipase A; PTEN, phosphatase and tensin homolog; TGFBR3, transforming growth factor beta receptor 3 \_\_\_\_\_ This is an open access article under the terms of the Creative Commons Attribution License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited.

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#### Abstract

As one of the most abundant immune cell populations in the tumor microenvironment (TME), tumor-associated macrophages (TAMs) play important roles in multiple solid malignancies, including breast cancer, prostate cancer, liver cancer, lung cancer, ovarian cancer, gastric cancer, pancreatic cancer, and colorectal cancer. TAMs could contribute to carcinogenesis, neoangiogenesis, immunesuppressive TME remodeling, cancer chemoresistance, recurrence, and metastasis. Therefore, reprogramming of the immune-suppressive TAMs by pharmacological approaches has attracted considerable research attention in recent years. In this review, the promising pharmaceutical targets, as well as the existing modulatory strategies of TAMs were summarized. The chemokine-chemokine receptor signaling, tyrosine kinase receptor signaling, metabolic signaling, and exosomal signaling have been highlighted in determining the biological functions of TAMs. Besides, both preclinical research and clinical trials have suggested the chemokine-chemokine receptor blockers, tyrosine kinase inhibitors, bisphosphonates, as well as the exosomal or nanoparticle-based targeting delivery systems as the promising pharmacological approaches for TAMs deletion or reprogramming. Lastly, the combined therapies of TAMs-targeting strategies with traditional treatments or immunotherapies as well as the exosome-like nanovesicles for cancer therapy are prospected.

#### **KEYWORDS**

chemokine-chemokine receptor, exosomeimmune suppression, metabolism, tumorassociated macrophages, tyrosine kinase receptor

## 1 | BACKGROUND

Cancer is a heterogeneous disease that is composed of numerous cell types, including both cancer and noncancer cells. Although macrophage content is highly variable in different tumor entities, macrophages are among the most prominent tumor-associated noncancer cell type in the tumor microenvironment (TME),<sup>1,2</sup> known as tumor-associated macrophages (TAMs). Current evidence suggests that TAMs engage in complex network interactions with cancer stem cells, cancer cells, endothelial cells, fibroblasts, T cells, B cells, and natural killer cells.<sup>3</sup> Their cross talks accelerate the formation of the immune-suppressive TME that not only stimulates cancer cell proliferation, neoangiogenesis, lymphangiogenesis, drug resistance, and distant metastasis, but also further recruits macrophages to establish a vicious feedback loop that continually reinforces the immune-suppressive TME. Recently, multiple signaling pathways have been identified as critical nodes mediating TAMs polarization and interactions with malignant cells. Small molecule inhibitors or monoclonal antibodies targeting TAMs signaling have

#### HIGHLIGHT

- The biofunctions and clinical implications of TAMs in cancer progression were summarized.
- The current pharmaceutical targets in TAMs regulation were introduced and discussed.
- The latest advancements of agents that were effective in TAMs depletion or reprogramming were summarized and discussed.
- The promising future of exosomal signaling for TAMs-targeted therapy was discussed and expected.

been demonstrated to effectively inhibit cancer development and metastasis. Therefore, TAMs have emerged as a central drug target for cancer therapy. Here, our review focuses on discussing the biological functions of TAMs, critical signaling and targets regulating TAMs activities, and current therapeutic strategies for treating malignancies that affect TAMs.

### 2 | MACROPHAGE POLARIZATION IN CANCER

Since the first description by llya llyich Mechnikov in 1882, phagocytes have been reported to be present throughout the body and to perform specific biological functions. For example, liver macrophages (Kupffer cells) eliminate pathogenic and waste products from circulation.<sup>4</sup> Brain-resident macrophages (microglia) contribute to the maintenance of engrams (i.e., the neural bases of memories)<sup>5</sup> by facilitating synaptic pruning. Langerhans cells in the skin are responsible for activating local inflammatory reactions and clearing pathogenic substances.<sup>6</sup> Phagocytes can also be infiltrated into the TME to perform pro- and/or antitumor functions. Usually, macrophages are considered as a plastic cell type capable of activating or polarizing into different statuses. Macrophages change their activation or polarization statuses in response to any potential entity, which is capable of being recognized by macrophages. The common stimuli including growth factors, cytokines/chemokines, hypoxia, exosomes, microbes, microbial products, nucleotide derivatives, antibody-Fc receptor stimulation, glucocorticoids, infection, and phagocytosis.<sup>7</sup> Therefore, it is hard to establish the common nomenclatures or standards for describing the activation or polarization properties of macrophages induced by various stimuli.<sup>7</sup> There are mainly four definitions of macrophage activation or

polarization statuses, including terms such as M1-like and M2-like, alternative and classical activation, "regulatory" macrophages, and subdivisions originating from the parent terms.<sup>7</sup> In 2012, Mills proposed the M1-like/M2-like dichotomy to describe the two major and opposing activities of macrophages.<sup>8</sup> At present, this definition method of macrophage activation or polarization statuses is widely used by the majority of researchers. M1-like polarization is reported to be induced by pathogen-associated patterns, including lipopolysaccharides (LPS), interferon- $\gamma$ , granulocyte-macrophage colony-stimulating factor (GM-CSF), and tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ). M1-like activation causes the release of various cytokines, such as interleukin 6 (IL-6), IL-1 $\beta$ , and TNF- $\alpha$ , that facilitate a proinflammatory response to defense against pathogenic insults.<sup>9</sup> In contrast, the M2-like phenotype is induced by damage-associated patterns (DAMPs) from cytokines such as IL-4 and IL-13, which subsequently activate the JAK-STAT pathway and turn on the expression of antiinflammatory cytokines, such as resistin-like molecule  $\alpha$ , IL-10, and arginase 1 (ARG1).<sup>10</sup> These signals further accelerate the remodeling of the TME to promote cancer angiogenesis, growth, and immune suppression. Usually, M1-like is considered to be an inhibitory phenotype due to its function in promoting Th1 responses that exert tumoricidal and microbicidal functions, whereas M2-like represents a restorative phenotype due to its effects in activating Th2 responses that contribute to tissue remodeling/repair, angiogenesis, immunosuppression, and tumor progression.<sup>11</sup> Generally, both M1-like and M2-like phenotypes can exist within the same microenvironment; therefore, the molecular targets controlling the polarization balance are considered as an important approach for cancer therapy. The polarization biomarkers of M1-like macrophages include CD86 and CD80, while the polarization biomarkers of M2-like macrophages include CD163, CD204, CD206, CD115, and CD301.<sup>12</sup> Particularly, CD163 can simultaneously be present both in a membrane-bound form on M2-like phenotype TAMs and a soluble form in plasma. Because of its macrophage-specific expression characteristic and highly elevated concentrations during various pathological conditions, soluble CD163 (sCD163) level is also regarded as a reliable biomarker for monitoring macrophage activity.<sup>13</sup> For example, CD163<sup>+</sup> TAMs were reported to express high levels of immune-checkpoint molecules PD1 and PD-L1. Blocking PD1/PD-L1 signaling in CD163<sup>+</sup> TAMs with antibodies could induce M1-like polarization and increase macrophage phagocytosis, reduce tumor burden and prolong survival,<sup>14,15</sup> and result in an increased release of sCD163 in the lesional skin of melanoma.<sup>13</sup> Meanwhile, Fujimura et al. identified serum sCD163 as a reliable predictive biomarker for immunerelated adverse events (irAEs) and efficacy of nivolumab

in patients with advanced melanoma.<sup>16,17</sup> These findings suggest that macrophage polarization modulation may also be the potential pharmacological mechanism of immune-checkpoint inhibitors. The combination treatment of macrophage-targeted strategy and PD-1/PD-L1 blockade may provide a synergetic antitumor efficacy, which deserves further investigations.

#### 3 | CLINICAL IMPLICATIONS OF TAMS IN CANCER DEVELOPMENT

Over the past decade, TAMs have attracted rising research attention for their modulatory potential on angiogenesis, cytokines, and immune regulation that either inhibit or facilitate tumor progression. Clinically, elevated infiltration levels of M1-like macrophages within tumor tissues predict a favorable prognosis, while elevations in the M2-like macrophages usually predict poor outcomes.<sup>18</sup> A tissue microarray analysis of 553 primary non-small cell lung cancer patients revealed that M1-like macrophages in metastatic lymph nodes were a predictor of improved survival.<sup>19</sup> In esophageal cancer patients, cases with high CD163 and CD204 expression levels showed a significantly shorter overall survival than those with comparatively lower CD163 and CD204 levels.<sup>20</sup> In addition, in triple-negative breast cancer patients with either high CD163 expression or low CD163 expression, the 5-year overall survival ratio was 72.3% or 82.5%, respectively.<sup>21</sup> These clinical observations have been well supported by experimental studies using macrophage depletion or overexpression strategies. For example, genetic ablation of GM-CSF in different murine tumor models-such as the MMTV-PyVT<sup>+/-</sup> breast cancer model, the spontaneous colon cancer model, and the osteosarcoma xenotransplant model—significantly reduces M2-like macrophage density in solid tumors, which in turn inhibits cancer growth and progression.<sup>22-24</sup> In addition, macrophage depletion with clodronate-encapsulated liposomes also yields a significant reduction of tumor volume in several malignancies.<sup>25,26</sup> In contrast, overexpression of CSF-1 remarkably increases TAMs recruitment and accelerates cancer development and metastasis.<sup>27</sup> It should be noted that the effect of TAMs on tumor tumorigenesis and progression may fluctuate due to their differentiation heterogeneity.<sup>28</sup> Based on these clinical observations and experimental findings, numerous studies have investigated the underlying mechanisms of TAMs in controlling cancer development. At present, M2-like macrophages have been found to participate in all steps of malignant development including carcinogenesis, neoangiogenesis, overall immune suppression, drug resistance, and later recurrence and/or metastasis (Figure 1).

### 3.1 | TAMs contribute to carcinogenesis and neoangiogenesis

TAMs have been found to be involved in the first step of carcinogenic lesion formation during neoplasia. Macrophage infiltration has been found to be upregulated in a murine chemically induced skin carcinogenesis model.<sup>29</sup> Similarly, a massive accumulation of CD206<sup>+</sup> or ARG1<sup>+</sup> macrophages has also been found in an inflammation-mediated skin tumorigenesis mice model, while macrophage ablation has been shown to significantly reduce tumor incidence.<sup>30</sup> In an EGFR-driven lung carcinogenesis model, sustained macrophage recruitment has been observed and macrophage depletion causes a significant reduction in tumor burden.<sup>31</sup> Neoangiogenesis is also a critical step during carcinogenesis, in which macrophage infiltration is also involved. Various studies have suggested that TAMs are predominantly located near the blood vessels of malignant solid tumors, and TAMs numbers are usually positively correlated with blood vessel density.<sup>32-35</sup> Functional studies have also demonstrated that TAMs elimination causes the reduction of neoangiogenesis,<sup>36</sup> while TAMs enhancement aggravates this process.<sup>36</sup> Mechanistic studies imply that TAMs can release multiple proangiogenic factors, such as vascular endothelial growth factor A (VEGF-A), macrophageinhibitory factor (MIF), adrenomedullin (ADM), plateletactivating factor (PAF), platelet-derived growth factor (PDGF), basic fibroblast growth factor (bFGF), and TGF- $\beta$ , as well as numerous cytokines such as TNF- $\alpha$ , IL-1, IL-8, and monocyte chemoattractant protein-1 (MCP-1).<sup>37-40</sup> Additionally, TAMs also release numerous angiogenesismodulating enzymes including iNOS,<sup>41</sup> COX-2, and matrix metalloproteinases (MMPs),<sup>42-44</sup> all of which have been associated in matrix degradation and endothelial cell invasion.

# 3.2 | TAMs facilitate the formation of the immune-suppressive microenvironment

TAMs recruitment not only supports cancer growth *via* neoangiogenesis induction but also facilitates the establishment of the immune-suppressive microenvironment. Recent studies have suggested that TAMs express PD-L1, PD-L2, CD86, and CD80, all of which induce CD8<sup>+</sup> T cell dysfunction upon binding to immune-checkpoint receptors such as PD1 or cytotoxic T-lymphocyte-associated protein 4 (CTLA4).<sup>44,45</sup> In addition, TAMs release multiple cytokines, enzymes, and chemokines that inhibit T-cell activity through natural regulatory T cell recruitment or L-arginine depletion in the TME. For example, IL-10 produced by TAMs could suppress IL-12



**FIGURE 1** Pro-oncogenic role of TAMs in cancer development. Macrophages could be differentiated into M1-like and M2-like phenotypes under different stimuli. M1-like macrophages were usually considered as a "killer" phenotype while M2-like macrophages were known as the "healing" phenotype. M2-like macrophages have been reported to participate in the process of carcinogenesis and neoangiogenesis *via* releasing proangiogenic factors, enzymes, and MMPs. Meanwhile, M2-like macrophages promoted cancer recurrence and metastasis *via* modulating epithelial–mesenchymal transition, extracellular matrix degradation, and facilitating premetastatic niche formation. On the other hand, M2-like macrophages could induce chemoresistance by exosomal signaling or cell–cell contact. Most importantly, M2-like macrophages contributed to establishing the immune suppression microenvironment by elevating the PD-1/CTLA4 signaling or inhibiting the bio-functions of cytotoxic T cells or dendritic cells

secretion from myeloid cells and promote Th2-type immune response.<sup>46</sup> The secretion of TGF- $\beta$  and PGE2 can impair the maturation process of dendritic cells, which subsequently compromise the balance between innate and adaptive immunity.<sup>47,48</sup> Immune-checkpoint

inhibitors have revealed successful therapeutic responses in multiple malignant tumors such as melanoma and lung cancers.<sup>49</sup> Unfortunately, only approximately 20% of cancer patients respond to immunotherapy, and mixed responses can limit therapeutic efficacies and lead to local recurrences and/or distant metastases.<sup>50</sup> Given the abundance and immune-suppressive properties of TAMs, targeting TAMs has been suggested as a promising approach to promote the efficacy of checkpoint antagonists. For example, anti-PD1/anti-CTLA4 treatment can decrease pancreatic tumor growth by approximately 50%, while their combination with PLX3397 (CSF1R inhibitor) can dramatically attenuate tumor expansion and even results in tumor regression by 15%.<sup>51</sup> Fc $\gamma$ R is a receptor typically expressed by TAMs. Similarly, a PD1 antibody also results in tumor growth inhibition in colon cancer xenografts, although this efficacy typically varies among mice. Strikingly, when a PD1 antibody and  $Fc\gamma R$  antagonist were simultaneously administrated, tumor growth was completely inhibited in all mice.<sup>52</sup> Additionally, suppressing IL-10 signaling in TAMs can promote CD8<sup>+</sup> T cellmediated antitumor immune responses in breast cancer receiving chemotherapy.<sup>46</sup> Given these encouraging preclinical results, utilization of TAMs-targeting strategies is promising for future clinical application as a combination therapy with checkpoint inhibitors.

# 3.3 | TAMs aggravate cancer drug resistance

TAMs density is also closely correlated with therapeutic responses and is highlighted in cancer drug resistance. Multiple studies have found that TAMs populations are enriched following toxic cancer-killing treatments. In aromatase-inhibitor-resistant breast cancer, TAMs density is much higher in primary tumor tissues and significantly correlates with decreased disease-free survival (DFS) and overall survival (OS).<sup>53</sup> TAMs infiltration was also found to be elevated following chemotherapy in a pancreatic ductal adenocarcinoma model, and TAMs pretreatment could render pancreatic cancer cells resistant to gemcitabine whereas this efficacy was strikingly enhanced by TAMs depletion *via* clodronate liposomes.<sup>54</sup> In colon cancer, TAMs infiltration has frequently detected been in chemoresistant patients and is significantly associated with poor survival.<sup>55</sup> The macrophage marker, CD68, is positively correlated with the expression of MDR1. Notably, blocking IL-6 secreted from TAMs is capable of resensitizing colon cancer cells to chemotherapeutic drugs. Meanwhile, B-cell activating factor released by TAMs is also effective in mediating bortezomib resistance in myeloma. In addition to cytokine secretion, ligandreceptor interactions between cells are also involved in mediating chemoresistance.<sup>56</sup> In a macrophage-myeloma contact model, plasma-membrane protein/gene profiling assays have demonstrated that CD18 and selectins in macrophages remarkably contribute to chemoresistance

*via* combining with intracellular cell adhesion molecule-1 (ICAM-1) and P-selectin glycoprotein ligand-1 (PSGL-1), respectively, in myeloma cells. Pharmacological inhibition or genetic knockdown of PSGL-1 or ICAM-1 signaling in these myeloma cells can diminish the protective effects of TAMs from cytotoxic cancer-killing agents including melphalan, doxorubicin, and bortezomib. Recent findings also suggest that exosomes are also implicated in TAMsinduced chemoresistance. It is found that macrophagesderived exosomal miR-21 signal could be directly delivered into gastric cancer cells, and thus results in apoptosis inhibition and cisplatin resistance of gastric cancer cells.<sup>57</sup> Therefore, it may be worthwhile to apply TAMstargeting therapies with traditional cancer-killing strategies in future clinical applications.

# 3.4 | TAMs accelerate cancer recurrence and metastasis

Local recurrence and distant metastasis are not solely determined by the malignant behavior of cancer cells because recent studies have suggested that stromal cells, particularly TAMs, act as an important driving force in these processes. For example, Giurisato et al. proved that extracellular-regulated protein kinase 5 (ERK5)-mediated macrophage proliferation could support melanoma invasiveness and metastasis in vivo, suggesting TAMs renewal as an integral component of tumor metastasis.<sup>58</sup> Mechanistically, numerous researches have demonstrated the crucial role of TAMs in regulating the epithelial-mesenchymal transition (EMT) process.<sup>59-61</sup> Cocultured hepatocellular cancer cells with TAMs could enhance the expression levels of mesenchymal markers including N-cadherin and vimentin, whereas attenuating the epithelial marker E-cadherin.<sup>60</sup> A similar phenomenon was also observed in breast, gastric, colon, and pancreatic malignancies.<sup>62</sup> Biologically, multiple TAMssecreted cytokines including IL-1 $\beta$ , IL-8, EGF, TNF- $\alpha$ , and TGF- $\beta$  have been validated to promote the EMT process.<sup>60,63,64</sup> Meanwhile, TAMs can secrete various kinds of proteolytic enzymes including MMPs, cathepsins, and serine proteases to degrade the extracellular matrix (ECM).<sup>65,66</sup> For instance, cathepsin B and S secreted by TAMs were found to be critical in promoting pancreatic cancer invasion.<sup>67</sup> An earlier study also demonstrated that TAMs synthesize SPARC/osteonectin to deposit collagen IV, resulting in enhanced tumor invasion and adhesion to other ECM components.<sup>68</sup> TAMs have also been shown to be necessary for facilitating cancer cell intravasation and extravasation. Multiphoton intravital imaging techniques have shown that intravasating cancer cells are invariably accompanied by a macrophage within one cell diameter.<sup>69</sup>

Interestingly, macrophages loss can significantly impair the extravasation rates and metastasis of cancer cells.<sup>70</sup> Additionally, TAMs have also been found to be critical in supporting the survival of cancer cells in the circulation. TAMs depletion via genetic methods can dramatically suppress cancer cell survival in pulmonary capillaries as well as the subsequent lung metastasis formation.<sup>71,72</sup> The underlying molecular mechanisms of their synergistic interaction are as follows. First, TAMs may trigger PI3K/Akt survival signaling in cancer cells to circumvent proapoptotic cytokines such as TRAIL.<sup>73</sup> Second, TAMs may alleviate survival stress from NK or cytotoxic T cells in the circulation.<sup>72</sup> Following extravasation. it has been found that TAMs are one of the key determinants in establishing the premetastatic niche (PMN).<sup>74</sup> Circulatory or residential TAMs can release chemokines that guide the localization of cancer cells into the PMN with increased expression levels of MMPs, fibronectins, S100A8, and S100A9.75-77 Our recent report demonstrated that TAMs-derived CXCL1 exerted an important role in recruiting breast cancer cells into the PMN.<sup>78</sup> Thus, TAMs act as an indispensable factor in fostering cancer cells traveling from the primary site to metastatic lesions. Molecular elucidation of TAMs regulation in cancer development is thus warranted and critical for further developing cancertargeting strategies.

# 4 | PHARMACEUTICAL TARGETS OF TAMS

To date, several key pathways including chemokine– chemokine receptor signaling, tyrosine kinase receptor (RTK) signaling, metabolic signaling, and exosomal signaling have been highlighted in determining the biofunctions of TAMs. The promising pharmacological targets of each signaling were described as following and summarized in Table 1.

# 4.1 | Chemokine-chemokine receptor signaling

Chemokines, which are small and soluble (8–14 kDa) signaling proteins, are a family of chemotactic cytokines responsible for cellular trafficking. More than 50 kinds of chemokines and 20 kinds of chemokine receptors have been identified until now.<sup>79</sup> According to the positions of conserved cysteine residues, chemokines can be classified into four groups including CXC, CC, CX3C, and C.<sup>80</sup> Correspondingly, chemokine receptor nomenclature essentially follows that of chemokines. Eleven kinds of CC chemokine receptors (CCR1–CCR11) recognize the

CC subfamily chemokines, while seven kinds of CXC receptors (CXCR1-CXCR7) recognize the CXC subfamily chemokines, respectively.<sup>80</sup> Similarly, the CX3C subfamily chemokine (CX3CL1) only binds to CX3CR1 receptor while the C subfamily chemokines (XCL1/2) only bind to XCR1 receptor, respectively. Chemokine receptors are typical Gprotein-coupled receptors (GPCRs) with seven transmembrane domains. Chemokine receptors can relay their signals through heterotrimeric G proteins that result in the directional migration of cells along a concentration gradient of ligands.<sup>79</sup> The interactions between chemokine and chemokine receptors are summarized in Figure 2 and Table 1. In the TME, TAMs are considered as the main stromal cells that secrete chemokines. As indicated by chemokine arrays, highly expressed chemokines of TAMs have been reported to consist of CCL2, CCL3, CCL5, CCL18, CXCL1, and CXCL12. Currently, the ligand receptor for CCL2 has been identified as CCR2. CCR1 and CCR5 have been found to be the receptors for both CCL3 and CCL5. Meanwhile, CCR3 has also been identified as a receptor for CCL5. By immunoprecipitation methods, the cognate receptor for CCL18 has been identified as PITPNM3, which has been suggested to be a putative sixtransmembrane protein that is sufficient to exert GPCRrelated functions. Additionally, the receptors for CXCL1 and CXCL12 have been identified to be CXCR2 and CXCR4, respectively.

Chemokines and chemokine receptors are important for attracting infiltrating leukocytes to the TME. CCL2 and CCL5 are the primary chemokines responsible for monocytic precursors recruitment in tumors.<sup>81-83</sup> Their expression levels are closely associated with TAMs infiltration, lymph-node metastasis, as well as unfavorable prognosis. CCL2 production from macrophages is critical for recruiting myeloid-derived suppressor cells (MDSCs) to build a local immunosuppressive microenvironment in gliomas.<sup>84</sup> Moreover, in mixed-bone marrow chimeric assays, CCR2-deficient MDSCs failed to accumulate in gliomas.<sup>84</sup> In addition, the CCL2-CCR2 axis was also found to be critical in mobilizing dendritic cell-like antigenpresenting cells into fibrosarcomas, as well as tumorinfiltrating lymphocytes.<sup>85</sup> Multiple studies have demonstrated that CCL5 can inhibit the antitumor immune response and is correlated with poor outcomes in multiple malignancies.<sup>86-89</sup> CCL5/CCR3 signaling has been found to promote Th2 immune polarization and results in luminal breast cancer.90 The CCL5/CCR5 chemokine axis is also capable of enhancing TGF- $\beta$ -mediated killing of cytotoxic CD8<sup>+</sup> T cells in colon cancer through regulatory T cells.<sup>91</sup> With regard to CXC chemokines, CXCL1 chemokines have been demonstrated to be effective in recruiting CXCR2<sup>+</sup> Treg cells and lead to metastasis in nonsmall cell lung cancer.92 CXCL1 is also critical in

TABLE 1 Th	e pharmaceutical tar	gets of TAMs		
Signaling	Targets	Tumors	Mechanisms	References
Chemokine– chemokine receptor signaling	CCL2/ CCR2	Breast cancer	Facilitating cancer progression via activation of ERα and PI3K/AKT/NF-κB signaling; inducing tamoxifen resistance by activating PI3K/Akt/mTOR signaling; enhancing growth and cell-cycle progression through SRC and PKC activation	177–179
	CCL2/ CCR2	Prostate cancer	Recruiting and regulating macrophages via the MAPK/ ERK signaling pathway; protecting cancer cells from autophagic death via the PI3K/Akt/survivin pathway; promoting cancer progression through the induction of MMP-2 activity	180–182
	CCL2/ CCR2	Liver cancer	Inducing invasion and EMT through activation of the Hedgehog pathway; inducing migration and invasion depending on MMP-2 and MMP-9;	183,184
	CCL2/ CCR2	Lung cancer	Increasing invasion via ERα; promoting TAMs infiltration via NF-κB/CCL2 signaling; promoting invasion mediated by autocrine loop of PDGF-BB	185–187
	CCL2/ CCR2	Ovarian cancer	Promoting peritoneal metastasis through P38-MAPK pathway	188
	CCL2/ CCR2	Gastric cancer	Inhibiting proapoptotic autophagy by activating PI3K-Akt-mTOR signaling	189
	CCL5/CCR1	Colorectal cancer	Promoting cancer progression through CCL5/β-catenin/Slug pathway	190
	CCL5/CCR1	Prostate cancer	Promoting invasion by increasing MMP-2/9 and activating ERK/Rac signaling	191
	CCL5/CCR5	Lung cancer	Increasing migration via PI3K/AKT/NF-κB signaling; Promoting metastasis and macrophages infiltration via EZH2	192,193
	CCL5/CCR5	Breast cancer	Promoting immune cell infiltration driven by TNFα/NF-κB signaling; enhancing Trastuzumab resistance by ERK pathway activation; promoting cancer proliferation through the mTOR pathway	194–196
	CCL5/CCR5	Ovarian cancer	Promoting invasion and migration via NF-κB–mediated MMP-9 upregulation; mediating differentiation of CSCs into ECs via the NF-κB and STAT3 signaling	197,198
	CCL5/CCR5	Liver cancer	Inducing EMT through activation of the Akt pathway	199
	CCL5/CCR5	Pancreatic cancer	Recruiting Treg cells via cancer-FOXP3; inducing proliferation of cancer cells through F-actin polymerization	200,201
	CCL5/CCR5	Colorectal cancer	Enhancing TGF-β-mediated killing of CD8 <sup>+</sup> T cells; Facilitating immune escape via the p65/STAT3-CSN5-PD-L1 pathway	91,202
	CCL5/CCR5	Gastric cancer	Enhancing aberrant DNA methylation via STAT3 signaling; inducing invasion and proliferation via KLF5	203,204
	CCL5/CCR5	Melanoma	Activating apoptotic pathway involving release of cytochrome c and activation of caspase-9 and caspase-3	205
	CCL18	Breast cancer	Inducing EMT via PI3K/Akt/GSK3β/Snail signaling through AnxA2; recruiting Treg cells and promoting metastasis and via PITPNM3; enhancing EMT via N-Ras/ERK/PI3K/NF-κB/Lin28b signaling	206–208

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TABLEI	(Continued)			
Signaling	Targets	Tumors	Mechanisms	References
	CCL18	Oral squamous cell carcinoma	Promoting growth and metastasis by activating the JAK2/STAT3 signaling; promoting migration via mTOR signaling through Slug	209,210
	CCL18	Ovarian cancer	Promoting migration through Pyk2 signaling or mTORC2 signaling	211,212
	CCL18	Lung cancer	Promoting migration and invasion via Nir1 through Nir1-ELMO1/DOC180 signaling	213
	CCL18	Bladder cancer	Promoting lymphangiogenesis by increasing the production of VEGF-C and MMP-2	214
	CCL18	Pancreatic cancer	Promoting progression and the Warburg effect via NF-κB/VCAM-1 pathway	215
	CCL18	Gastric cancer	Promoting invasion and migration via ERK1/2/NF-κB signaling	216
	CCL20/CCR6	Lung cancer	Promoting migration via ERK1/2-MAPK and PI3K pathways; promoting progression mediated by lncRNA-u50535 through CCL20/ERK signaling	217,218
	CCL20/CCR6	Renal cell carcinoma	Inducing EMT through Akt activation	219
	CCL20/CCR6	Colorectal cancer	Promoting chemoresistance via FOXO1/CEBPB/NF-κB signaling; promoting growth and metastasis mediated by lncRNA-u50535; promoting proliferation and migration through ERK-1/2, SAPK/JNK, and Akt signaling	220-222
	CCL20/CCR6	Pancreatic cancer	Increasing TRAIL resistance via RelA-CCL20 pathway; promoting migration, EMT, and invasion through PI3K/AKT-ERK1/2 signaling	223,224
	CCL20/CCR6	Breast cancer	Increasing chemoresistance via ABCB1/ NF- $\kappa$ B signaling; promoting bone metastasis mediated by HuR; inducing EMT via PKC- $\alpha$ /Src/Akt/NF-kB/Snail signaling; promoting invasion by PKC- $\alpha$ through EGFR and ERK1/2/MAPK pathway	225-228
	CCL20/CCR6	Gastric cancer	Inducing EMT mediated by CRKL via Akt pathway	229
	CCL22/CCR4	Prostate cancer	Promoting migration and invasion via phosphorylation of Akt	230
	CCL22/CCR4	Colorectal cancer	Enhancing chemoresistance via PI3K/AKT pathway and caspase-mediated apoptosis	231
	CCL22/CCR4	Liver cancer	Recruiting regulatory T cells through p65/miR-23a/CCL22 axis	232
	CXCL1/CXCR2	Lung cancer	Recruiting Treg cells via miR141-CXCL1-CXCR2 signaling	92
	CXCL1/CXCR2	Breast cancer	Promoting metastasis via activating NF- $\kappa$ B/SOX4 signaling; promoting cancer cell survival mediated by TNF- $\alpha$ via NF- $\kappa$ B signaling; promoting metastasis and chemoresistance through myeloid cell-derived S100A8/9; stimulating migration and invasion via activation of the ERK/MMP2/9 signaling	105,233-235
	CXCL1/CXCR2	Pancreatic cancer	Promoting migration and invasion by CXCL1-mediated Akt phosphorylation	236
	CXCL1/CXCR2	Ovarian cancer	Recruiting MDSC mediated by Snail; Enhancing metastatic potential via the TAK1/NF-κB signaling; driving cancer progression by NF-κB activation via EGFR-transactivated Akt signaling	237-239

TABLE 1	(Continued)			
Signaling	Targets	Tumors	Mechanisms	References
	CXCL1/CXCR2	Prostate cancer	Promoting migration via the Src activation; increasing migration and invasion by fibulin-1 downregulation through NF-κB/HDAC1	240,241
	CXCL1/CXCR2	Colorectal cancer	Forming a premetastatic niche stimulated by VEGFA	93
	CXCL1/CXCR2	Gastric cancer	Promoting migration and metastasis through activation of CXCR2/STAT3 signaling; promoting lymph node metastasis through integrin $\beta$ 1/FAK/AKT signaling; promoting tumor growth through VEGF pathway activation	242-244
	CXCL12/CXCR4	Colorectal cancer	Promoting proliferation, invasion, angiogenesis via MAPK/PI3K/AP-1 signaling; enhancing metastatic potential via PI3K/Akt/mTOR pathway; promoting progression by lncRNA XIST/ miR-133a-3p/ RhoA signaling; promoting growth and metastasis through activation of the Wnt/β-catenin pathway; promoting chemoresistance via surviving	245-249
	CXCL12/CXCR4	Breast cancer	Promoting metastasis by Pit-1-CXCL12-CXCR4 axis; driving the metastatic phenotype through activation of MEK/PI3K pathway	250,251
	CXCL12/CXCR4	Ovarian cancer	Recruiting MDSCs mediated by PGE2; promoting invasion through suppressing ARHGAP10 expression; promoting growth and migration through Notch pathway	94,252,253
	CXCL12/CXCR4	Lung cancer	Suppressing cisplatin-induced apoptosis through JAK2/STAT3 signaling	254
	CXCL12/CXCR4	Prostate cancer	Inducing myofibroblast phenoconversion through EGFR/MEK/ERK signaling; promoting migration and invasion through SLUG	255,256
	CXCL12/CXCR4	Gastric cancer	Inducing migration via SRC-mediated CXCR4-EGFR; increasing invasiveness via integrin β1 clustering; promoting migration by F-actin reorganization and RhoA activation through mTOR signaling	257-259
	CXCL12/CXCR4	Thyroid papillary cancer	Enhancing proliferation and invasion through Akt and snail signaling	260
	CXCL8/ CXCR1/2	Gastric cancer	Inhibiting CD8+ T cells function by inducing the expression of PD-L1	261
	CXCL8/ CXCR2	Lung cancer	Stimulating cell proliferation via transactivation of the EGFR involving the MAPK pathways	262
	CXCL8/ CXCR1/2	Breast cancer	Increases the activity of cancer stem-like cells by transactivation of HER2	263
	CXCL8/ CXCR1/2	Melanoma	Increasing tumor growth and metastasis by activating MMP-2	264
	CXCL8/ CXCR1/2	Prostate cancer	Promoted tumorigenesis via STAT3/MALAT1 pathway	265
	CXCL8/ CXCR1/2	Colorectal cancer	Enhancing resistance of cancer cells to anoikis through AKT/TOPK/ERK signaling; enhancing migration by increasing $\alpha v \beta 6$ integrin expression; promoting proliferation and migration through heparin binding EGF	266-268
	CX3CL1/CX3CR1	Breast cancer	Promoting the migration and invasion via Src/FAK signaling; triggering proliferation through transactivation of EGF pathway	269,270

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# TABLE 1 (Continued)

CLINICAL AND TRANSLATIONAL MEDICINE

TABLET (C	continueu)			
Signaling	Targets	Tumors	Mechanisms	References
	CX3CL1/CX3CR1	Lung cancer	Promoting the migration and invasion via Src/FAK signaling; promoting lymph node metastasis via JNK and MMP2/MMP9 pathway	271,272
	CX3CL1/CX3CR1	Ovarian cancer	Promoting EMT mediated by HIF-1 $\alpha$	273
	CX3CL1/CX3CR1	Prostate cancer	Promoting EMT via TACE/TGF-α/EGFR pathway and upregulation of Slug; promoting metastasis via EGFR through activation of the Src/FAK pathway	274,275
	CX3CL1/CX3CR1	Pancreatic cancer	Enhancing growth and migration through JAK/STAT signaling; promoting motility, invasion, and growth via AKT activation; promoting tumor cell survival and TRAIL resistance via RelA/NF- <i>κ</i> B signaling; stimulating HIF-1 <i>α</i> expression through the PI3K/Akt and MAPK pathways	276–279
Tyrosine kinase receptor signaling	CSF-1/CSF-1R	Gliomas	Promoting tumor cell survival through IGF-1R/ PI3K signaling	280
	CSF-1/CSF-1R	Breast cancer	Driving tumor progression through Kindlin-2/TGFβ/CSF-1 signaling	281
	CSF-1/CSF-1R	Ovarian cancer	Promoting metastasis through the induction of MMP-9 activity	282
	CSF-1/CSF-1R	Pancreatic cancer	Remodeling tumor immune microenvironment via PI3K-γ and CSF-1/CSF-1R pathways	283
	Ron	Prostate cancer	Promoting prostate cancer cell growth via MST1R	284
	Ron	Pancreatic cancer	Promoting macrophage polarization through MST1-MST1R signaling	285
Metabolic signaling	Glycolysis	Pancreatic cancer	Conferring a prometastatic phenotype by HK2;	286
	Glycolysis	Breast cancer	Promoting M2-TAMs polarization through activation of GPR132 by lactate; promoting tumor angiogenesis by hypoxia-induced REDD1/ mTOR signaling	287,288
	Glycolysis	Melanoma	Promoting M2-TAMs accumulation by HMGB1; inducing TME acidification involving GPCR	141,289
	Glycolysis	Thyroid carcinoma	Inducing reprogramming of TAMs and inflammation through AKT1/mTOR pathway	140
	Glycolysis	Lung cancer	Increasing glucose uptake and glycolysis flux by activation of AMPK; promoting M2-TAMs polarization mediated by HIF1α	290,291
	Fatty acid metabolism	Bladder carcinoma	Promoting cancer growth and metastasis through COX-2/PGE2 pathway	292
	Fatty acid metabolism	Liver cancer	Promoting cell migration by enhancing IL-1β secretion; enhancing the accumulation and polarization of M2-like TAMs by RIPK3	293,294
	Fatty acid metabolism	Breast cancer	Promoting cancer progression by FABP by favoring IL6 / STAT3 signaling; promoting TAMs differentiation through caspase-1/PPARγ/MCAD pathway	295,296
	Fatty acid metabolism	Ovarian cancer	Promoting the M2-like polarization through the PPARγ/NF-κB pathway	297
	Glutamine synthesis	Lung cancer	Promoting M2-like polarization through GS	298

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IABLE I	(Continued)			
Signaling	Targets	Tumors	Mechanisms	References
	Tryptophan metabolism	Colon, gastrointestinal cancer	Inhibiting T-cell functions through IDO upregulation	299,300
	Arginine metabolism	Melanoma	Promoting tumor cell proliferation through ARG1	301
	Arginine metabolism	Breast cancer	Promoting cell growth by ARG induction and suppressing NO-mediated tumor cytotoxicity	302
	Iron metabolism	Breast cancer	Releasing iron mediated by FPN in the TME; supporting proliferation by LCN	303,304
	RAGE	Breast cancer	Enhancing tumor growth and metastasis through S100A7 signaling	305
	RAGE	Glioma	Promoting angiogenesis through MMP9 signaling	306
	RAGE	Melanoma	Promoting invasiveness through S100A4 signaling	307
Exosomal signaling	miR-21-5-p, miR-155-5p	Colorectal cancer	Promoting migration and invasion by downregulating expression of BRG1	164
	miR-25-3p, miR-130b-3p, miR-425-5p	Colorectal cancer	Inducing M2-like macrophage polarization by regulating PTEN/PI3K/Akt signaling	308
	Wnt	Colorectal cancer	Promoting chemoresistance through Wnt signaling	309
	miR-125b-5p	Melanoma	Educating TAMs by targeting LIPA	310
	miR-21	Gastric cancer	Suppressing apoptosis through PI3K/AKT signaling by down-regulation of PTEN	57
	miR-301a	Pancreatic cancer	Inducing M2-like macrophage polarization via PTEN/PI3Kγ signaling	311
	miRNA-501-3p	Pancreatic cancer	Promoting progression through the TGFBR3-mediated TGF- $\beta$ signaling	312
	miR-223	Ovarian cancer	Promoting drug resistance via the PTEN-PI3K/AKT pathway	313
	miR-95	Prostate cancer	Promote cell proliferation, invasion, and EMT via miR-95/JunB axis	314

facilitating PMN formation in colorectal cancer by recruiting CXCR2<sup>+</sup> MDSCs.<sup>93</sup> Meanwhile, CXCL12/CXCR4 signaling has been reported to induce the recruitment of MDSCs in the TME of human ovarian cancer.94 Besides shaping the tumor immune milieu, chemokine axis is also critical in regulating the neoangiogenic process. CXC chemokines have been reported to be directly chemotactic for endothelial cells and to stimulate angiogenesis in vivo.<sup>95</sup> CXCR4 receptor is expressed on endothelial cells and its activation after CXCL12 binding can induce the proliferation and migration of endothelial cells. Moreover, CXCL12 recruits CXCR4<sup>+</sup> circulating or bonemarrow endothelial precursors and therefore promotes tumor angiogenesis.<sup>96</sup> Similarly, CXCL1 was also reported to promote neoangiogenesis in breast, liver, and colorectal cancers.<sup>97–99</sup> Finally, chemokine signaling can directly act on chemokine receptors expressed on the plasma membrane of cancer cells, thus controlling their malignancyrelated functions. For example, a variety of chemokine

receptors—including CXCR2, CXCR4, CCR1, CCR2, and CCR5—have been found in breast tumor cells.<sup>100–104</sup> It is found that CXCL1 can induce chemoresistance and metastasis of breast cancer *via* CXCR2 activation.<sup>105</sup> Furthermore, CCL2 induces hepatocellular carcinoma invasion and EMT *via* CCR2 activation.<sup>106</sup> CXCL12 is also capable of regulating glioblastoma-stem-like cells by modulating CXCR4.<sup>107</sup> Moreover, CXCL12-CXCR4 signaling has also been highlighted in guiding the homing of cancer cells to their specific metastatic organs.<sup>108</sup> Altogether, these diverse chemokine functions establish crosstalk between cancer cells and the TME, which represents a promising target for the successful development of novel therapeutic strategies.

It should be pointed out that multiple chemokines play multifaceted roles within the complicated TME, usually dividing into two directions: paracrine signaling for immune activation and autocrine signaling for proliferation and metastasis of cancer cells.<sup>109–111</sup> For example, the



**FIGURE 2** Pharmaceutical targets involved in TAMs regulation. Chemokine–chemokine receptor signaling was the most important target for TAMs-targeting therapy, particularly for CCL2-CCR2, CCL5-CCR5, CXCL1/CXCR2, and CXCL12/CXCR4 signaling. Tyrosine kinase receptors including CSF-1R and RON were also highly expressed on the cell membrane of TAMs and perfect for drug intervention. Molecular targets involved in glycolysis, fatty acid oxidation, and amino acid metabolism in TAMs were also being focused on drug development. Notably, exosomes act as the ideal drug-loading and delivery vehicle for modulating biofunctions of TAMs

CXCL9-10 axis could induce the migration, differentiation, and activation of the CXCR3<sup>+</sup> activated T or NK cells, leading to increased antitumor immune response.<sup>112</sup> Additionally, CCL5/CCR5 axis is also a double-edged sword in cancer. On one hand, CCL5 could promote CSCs self-renewal and recruit the immunosuppressive cells (such as CCR5<sup>+</sup> TAMs and Tregs) to form a tumor-protecting TME. On the other hand, CCL5 could also promote antitumor immunity by recruiting CCR5<sup>+</sup> activated T cells and DCs to the TME and therefore enhances the immunotherapy response in different tumor types.<sup>110,111</sup>

TAMs-related chemokine signals have been widely investigated in skin area because of their reliable prognosis and therapeutic values for skin tumors. For example, Wu et al. reported that TAMs-related chemokine signals played key roles in the development and progression of cutaneous T-cell lymphoma (CTCL). Compared with normal controls, mycosis fungoides (the most common variant of CTCL) skin displayed abundant TAMs infiltration as well as increased expression of a subset of macrophagerelated chemokines and associated receptors.<sup>113</sup> Selective depletion of M2-phenotype TAMs using clodronatecontaining liposomes remarkably delayed mycosis fungoides development in vivo. Interferons (IFNs) have been widely used for mycosis fungoides and malignant melanoma treatment in the clinic. Growing evidence has suggested that modulating TAMs-related chemokine signals may be the possible mechanism of the therapeutic effects of IFNs. For example, Furudate et al. reported that IFN- $\alpha$ 2a and IFN- $\gamma$  stimulation could decrease the release of CCL17/18, whereas elevating the production of CXCL10/11 in monocyte-derived TAMs in vitro. More importantly, the subcutaneous administration of IFN- $\alpha$ 2a could increase the CXCL11-secreting cell population in the lesional skin of advanced mycosis fungoides patients.<sup>114</sup> Kakizaki et al. reported that peritumoral injection of IFN- $\beta$  remarkably suppressed the recruitment of PD-L1<sup>+</sup> TAMs in B16F10 melanoma xenograft in vivo.<sup>115</sup> Meanwhile, IFN- $\beta$  administration dramatically attenuated the production of CCL17/22, whereas elevated the secretion of CXCL9/11 from TAMs. More importantly, these immunomodulatory effects of IFN- $\beta$  on TAMs were also observed in the lesional skin of patients with in-transit melanoma. It should be noted that the clinical application of IFNs for the treatment of cancer is still controversial. For example, Numerous clinical trials have examined the antitumor potential of recombinant IFN- $\gamma$  as an adjuvant to surgery or conventional chemotherapy.<sup>116</sup> However, these clinical trials often reported conflicting outcomes in patients with the same tumor type.<sup>116</sup> Extensive efforts are still needed to better understand the complex roles of IFNs in cancer treatment.

## 4.2 | RTK signaling

RTKs play a vital role in signal transduction that supports cellular communication and survival. Dysregulation of RTKs, such as receptor mutation or overexpression, contributes to the initiation, tumorigenesis, and progression of multiple malignancies. To date, it has been reported that 58 kinds of RTKs are edited by the human genome.<sup>117,118</sup> Notably, colony stimulation factor-1 receptor (CSF-1R), one of the most important members of the RTKs family, is highly expressed by TAMs.<sup>119</sup> Upon binding to its ligands including CSF-1 or IL-34, CSF-1R undergoes dimerization and therefore causes a signaling cascade that facilitates the proliferation, functioning, and survival of macrophages. High expression of CSF-1 has been found in breast, prostate, pancreatic, gastric, and many other types of cancers, and its level has been closely associated with TAMs density and poor prognosis.<sup>120-123</sup> It is found that CSF-1-positive cancer cells colocalize with abundant CSF-1R-positive TAMs in invasive breast cancer, and the CSF-1/CSF-1R axis is significantly correlated with a higher grade, basal-like phenotype, and lymph node metastasis.<sup>124</sup> In CSF-1-knockout MMTV-PyMT<sup>+/-</sup> spontaneous breast cancer mice, the perivascular macrophage density was declined with a sixfold reduction, accompanied by a 16-fold decrease of circulating tumor cells in the blood.<sup>125</sup> Moreover, CSF-1/CSF-1R signaling of macrophages could suppress CD4<sup>+</sup> T-cell responses and induce the generation of Treg cells, which facilitated the immune-suppressive TME.<sup>126</sup> Additionally, CSF-1 is also involved in the proangiogenic process since it promotes M2-like polarization of macrophages and secretes a variety of angiogenic factors.<sup>127</sup> Therefore, blocking CSF-1/CSF-1R activation is becoming an important therapeutic strategy to inhibit cancer growth and metastasis.

RON is another RTK expressed on macrophages surface that specifically recognizes macrophage-stimulating protein (MSP). Overexpression of RON has been implicated in the progression and metastasis of diverse malignancies. In breast cancer, MSP/RON overactivation is sufficient to promote cancer growth as well as metastasis to the lungs, liver, brain, and bone. Conversely, RON inactivation is capable of inhibiting the growth, angiogenesis, and metastasis of breast cancer xenografts.<sup>128</sup> The cancer-promoting effects of MSP/RON signaling are mainly attributed to the M2-like phenotype polarization of macrophages. RON activation can cause the phosphorylation of STAT3, which is essential for the tumor-promoting efficacies and immunosuppressive functions of TAMs.<sup>129</sup> By contrast, the MSP/RON pathway can downregulate STAT1 activity, which actively participates in antitumor immune responses via up-regulating IL-12.<sup>130</sup> Eyob et al. has demonstrated that RON signaling in macrophages is capable of impairing antitumor functions of CD8<sup>+</sup> T cells.<sup>131</sup> Altogether, both STAT3 activation and STAT1 inactivation induced by MSP/RON signaling may contribute to tumor immune tolerance and lead to cancer progression. Considering the important role of RON receptor signaling in TAMs modulation, pharmacological inhibition of RON kinases has received more attention from research groups around the world.

Mer tyrosine kinase (MerTK) is also a macrophagespecific RTK and its overexpression usually correlates with poor prognosis in cancer.<sup>132</sup> MerTK signaling activation can promote tumor immune tolerance by initiating macrophages efferocytosis and promoting the clearance of immunogenic antigens. Meanwhile, MerTK signaling activation can suppress inflammatory cytokines production and polarize macrophages toward the anti-inflammatory M2-like phenotype.<sup>133</sup> In contrast, macrophages MerTK signaling blockage can induce the repolarization of macrophages toward the M1-like phenotype and enhance antitumor immunogenic cell death, a crucial step in Tcell priming and activation.<sup>134</sup> Thus, MerTK blockade may represent a promising antitumor strategy that could significantly increase tumor immunogenicity and potentiate antitumor immunity. It has been reported that MerTK knockdown or MerTK neutralizing antibody treatment could remodel the cellular immune profile, bringing a more inflamed TME with increased T cell infiltration and T cell-mediated cytotoxicity.<sup>132</sup> Additionally, anti-MerTK antibody treatment could stimulate T cell activation and synergize with anti-PD-1 or anti-PD-L1 therapies in tumorbearing mice.<sup>135</sup> Altogether, macrophages MerTK blockade could promote tumor immunogenicity and potentiate antitumor immunity, which represents a promising strategy for tumor immunotherapy.

#### 4.3 | Metabolic signaling

A typical feature of cancer cells is their abnormal metabolism caused by oxygen deprivation and nutrient limitation.<sup>136</sup> Hypoxic TME not only induces the metabolism pattern shifting of cancer cells but also reprograms the polarization balance and metabolic mode of macrophages, which facilitate the formation of immunesuppressive TME. In hypoxic regions of tumors, a more dominant M2-like phenotype is observed, which might indicate the potential effects of hypoxia on macrophage polarization.<sup>137</sup> Hypoxia-inducible factors (HIFs) are major transcriptional factors that are responsive to hypoxia and control aerobic glycolysis.<sup>138</sup> Enhanced glycolytic activity allows macrophages to synthesize sufficient ATP and various biosynthetic intermediates to maintain their tumor-prone functions. A recent study demonstrated the enhanced glycolytic activity of TAMs when compared with that of bone marrow-derived macrophages (BMDMs). Glycolytic proteins, such as hexokinase-2, phosphofructokinase, and enolase 1, were all increased in TAMs.<sup>139</sup> Similar findings were also observed in a transwell co-culture model of macrophages with cancer cells or tumor-extract solutions.<sup>140</sup> In addition to intrinsic HIF activation, cytokines and lactate released from cancer cells under hypoxic conditions also contribute to the glycolytic phenotype of TAMs. In hypoxic melanoma cells, HIF-1 $\alpha$ accumulation induces HMGB1 translocation that causes IL-10 production, which consequently promotes M2-like macrophage activation.<sup>141</sup> Cancer-cell-derived lactate has also been reported to reprogram macrophages to a tumor-prone phenotype by inhibiting the NF- $\kappa$ B pathway

that in turn may impair M1-like activation and suppress the immune surveillance functions of infiltrating T cells and NK cells.<sup>142–144</sup> Interestingly, the OXPHOS chain has also been found to be elevated in a coculture model of thyroid-carcinoma cells and macrophages, characterized by enhanced oxygen consumption rate.<sup>140</sup> The concentrations of AcCoA and succinate are significantly increased in the mitochondria of M2-like macrophages, suggesting that TAMs have an intact OXPHOS despite an impaired TCA cycle.<sup>145</sup> The enhanced OXPHOS chain may not only yield ROS-induced DNA damage but may also provide citrates for fatty acid synthesis. However, more studies are still required to reveal the precise switching mechanisms between glycolysis and OXPHOS in TAMs depending on cancer types and stages.

Fatty acid metabolism is important in cancer development. It has been suggested that increased fatty acid metabolism provides cancer cells with sufficient membranes and energy to support cancer growth.<sup>146</sup> It is found that fatty acid oxidation is necessary for IL-4-induced macrophage activation, as CPT1 suppression could attenuate the expression of the M2-like marker proteins, CD301, and CD206.147 Meanwhile, fatty acid oxidation can activate the downstream COX2/PGE2 pathway to promote cancer growth and metastasis. TAMs express high levels of fatty acid synthase and increased PPAR signaling, which promote fatty acid oxidation and tumor growth.<sup>148</sup> PPARs are composed of three isoforms including PPAR- $\alpha$ , PPAR- $\beta/\delta$ , and PPAR- $\gamma$ . PPAR- $\alpha$  manages lipid oxidation and clearance. PPAR- $\gamma$  facilitates lipogenesis while PPAR- $\beta/\delta$  controls lipid uptake and storage.<sup>149</sup> PPAR- $\beta/\delta$  plays a crucial role in mediating M2-like polarization.<sup>150</sup> Ovarian cancer patient-derived TAMs exhibited increased expression levels of multiple PPAR- $\beta/\delta$  target genes when compared with that of monocyte-derived macrophages.<sup>151</sup> Lipidomic analysis revealed that the deregulation of PPAR- $\beta/\delta$  target genes did not result from PPAR- $\beta/\delta$  overexpression or its promotion effect on target genes, but was attributed to high levels of polyunsaturated fatty acids, especially arachidonic acid and linoleic acid in tumor ascites.<sup>151</sup> The internalization of polyunsaturated fatty acids and the subsequent lipid droplet formation may provide a reservoir of PPAR- $\beta/\delta$  ligands to TAMs, leading to a consistent activation of PPAR- $\beta/\delta$  target genes, such as pyruvate dehydrogenase kinase 4 (PDK4). Interestingly, PDK4 can induce glucose catabolism shifting toward aerobic glycolysis, further rendering TAMs to adapt to hypoxic TME.<sup>151</sup> PPAR- $\gamma$  is also known to regulate metabolic pathways and M2-like polarization. Macrophage deficiencies in PPAR- $\gamma$  and  $\delta$  and their coactivator, PGC1*β*, are ineffective for IL-4-induced fatty acid oxidation and M2-like polarization.<sup>152</sup> Hence, targeting of PPAR signaling is a promising approach to control macrophage activation and cancer progression.

In addition to enhanced glycolysis and fatty acid oxidation, M2-like macrophages have also been found to increase glutamine synthesis through glutamate, which has not been demonstrated in M1-like macrophages.<sup>153</sup> Inhibition of glutamine syntheses results in a 50% reduction of M2-like polarization, which might be ascribed to the role of glutamine in protein glycosylation.<sup>154</sup> Meanwhile, glutamine is extensively fluxed into the TCA cycle to provide enough carbons to satisfy energetic needs.<sup>155</sup> Besides glutamate, another amino acid, arginine, is also required in the ARG1 pathway of M2-like macrophages.<sup>156</sup> Interestingly, ARG1 is also the downstream gene of HIF- $1\alpha$ . ARG1 overexpression not only enables cancer cells to rapidly adapt to the hypoxic microenvironment but also results in the production of ornithine and urea.<sup>156</sup> Ornithine can be further utilized in the synthesis of proline and polyamine, which are required for cellular proliferation, tissue remodeling, and collagen biogenesis.<sup>157</sup> Moreover, the arginine clearance by ARG1 suppresses Tcell immune responses, favoring the establishment of the immunosuppression microenvironment.<sup>157</sup> Besides, trvptophan metabolism has been shown to be deregulated in cancers. It was found that the enzyme accounting for tryptophan catabolism, indoleamine-2, 3-dioxygenase (IDO), was frequently overexpressed in M2-like macrophages.<sup>158</sup> IDO upregulation was reported to inhibit T-cell functions, whereas tryptophan treatment or administration of IDO inhibitors could restore T-cell proliferation.<sup>158</sup>

RAGE (receptor for advanced glycation end-product) is a member of the immunoglobulin superfamily and has been reported to be highly expressed on the surface of macrophages.<sup>159</sup> It has been reported that RAGE overexpression could induce the accumulation of TAMs and thus accelerate the growth of lung cancer xenografts in vivo.<sup>160</sup> Mechanistically, RAGE expression strictly correlates to the metabolic switch in cancer and immune cells. For example, cancer cells usually exhibit a metabolism switch characterized by decreased mitochondrial oxidative phosphorylation as well as elevated glycolysis, leading to the overproduction of toxic glucose metabolites.<sup>161</sup> The toxic glucose metabolites such as lactate, sorbitol, diacylglycerol (DAG), and methylglyoxal (MG) could further contribute to the development of advanced glycation end products (AGEs). On one hand, the formation of AGEs could impair the phagocytosis function of M1-like macrophages within the TME.<sup>162</sup> On the other hand, AGEs could induce RAGE expression in macrophages and their interaction could further promote cancer progression and metastasis by recruiting TAMs.<sup>160,163</sup> Therefore, AGEs/RAGE signaling may also represent a metabolism-related target for the pharmacological modulation of TAMs, which still needs further investigations. Collectively, the above findings suggest that the metabolic pathways and targets can also be applied

to modulate the biological functions of TAMs, which may provide valuable opportunities for treating multiple malignancies (Figure 2 and Table 1).

### 4.4 | Exosomal signaling

In recent years, exosomal signaling has evoked increased interest in cancer research. Exosomes are defined as small extracellular vesicles that are 30-100 nm in diameter.<sup>164</sup> Similar to cells, exosomes are composed of a lipid bilayer containing all known molecular components including DNA, RNA, and proteins.<sup>165</sup> Exosomes can be released by all cell types and widely exist in human bodily fluids such as blood, urine, amniotic fluid, bronchoalveolar lavage fluid, and malignant fluid. In particular, normal human blood is expected to contain approximately 2,000 trillion exosomes, while this value is elevated to 4,000 trillion in the blood of cancer patients.<sup>166–168</sup> Therefore, exosomal signaling is highly implicated in intercellular communication and carcinogenesis. Recent studies have also demonstrated that exosomal signaling plays a vital role in tumor angiogenesis,<sup>169</sup> metastasis,<sup>170</sup> drug resistance,<sup>171</sup> and immune regulation,<sup>172</sup> particularly for exosomes derived from macrophages. It has been reported that TAMs-derived exosomes exhibit high expression levels of miR-21-5-p and miR-155-5p, which significantly promote cellular migration and invasion of colorectal cancer cells.<sup>164</sup> Besides, exosomes from macrophages also contain a high level of Wnt, which is considered to provide critical signaling in EMT induction and mesenchymalphenotype maintenance.<sup>173,174</sup> In addition to metastasis, TAMs-derived exosomes also contribute to extracellularmatrix remodeling. It has been shown that MMP-12 and MMP-13, as well as cathepsin B, D, K, L, S, and Z, are all upregulated in TAMs-derived exosomes, suggesting their positive roles in extracellular remodeling.<sup>175</sup> Notably, exosomes released from TAMs also contain miRNAs that lead to Treg/Th17 cell imbalance in ovarian cancer, resulting in the generation of an immune-suppressive TME and thus promoting cancer progression and metastasis.<sup>176</sup> Altogether, targeting TAMs-derived exosomal signaling represents a novel and promising approach for cancer treatment (Table 1).

## 5 | PHARMACOLOGICAL MODULATION OF TAMS

### 5.1 | Agents targeting chemokine-chemokine receptors

Chemokine-chemokine receptor signaling constitutes an important network that regulates cancer growth, angiogenesis, immune suppression, and metastasis. Targeting of chemokine-chemokine receptor networks has been evaluated in multiple preclinical models and clinical trials, particularly in terms of the CCL2-CCR2, CCL5-CCR5, and CXCL12-CXCR4 axes (Table 2). Blockade of CCL2 with a neutralizing antibody inhibited inflammatory monocyte recruitment, reduced lung metastasis, and prolonged OS in breast cancer-bearing mice.<sup>315</sup> In a metastatic prostate cancer model, combined treatment with an anti-CCL2 antibody and docetaxel caused reduced tumor burden and bone resorption, as well as an improved survival period.<sup>316</sup> The development of CCR2 inhibitors has also yielded encouraging results in cancer treatments. Oral administration of the CCR2 inhibitor, PF-04136309, in a preclinical pancreatic cancer model reduced the number of tumor-infiltrating macrophages and inflammatory monocytes.<sup>317</sup> Significantly, PF-04136309 acted synergistically with the chemotherapeutic drug, gemcitabine, to inhibit metastasis, increase anti-tumor T-cell responses, and ultimately lead to an enhanced chemotherapeutic efficacy.<sup>318</sup> Similarly, the synergistic effects between PF-04136309 and paclitaxel or the standard chemotherapy, FOLFIRINOX, have also been reported in pancreatic cancer patients.<sup>319</sup> Another CCR2 inhibitor, CCX872, was demonstrated to improve anti-PD-1 treatment in preclinical settings and positive results were also obtained when used in combination with the FOLFIRINOX strategy in pancreatic cancer patients.<sup>320</sup> In murine models of hepatocellular carcinoma, combined treatment with the CCR2 antagonists, RDC018 or 747, with sorafenib was validated to inhibit tumor growth and metastasis, accompanied by a significant reduction of macrophage infiltration.<sup>321,322</sup> It should be pointed out that there are also some disappointing results in the clinical trials of CCL2/CCR2 blockers. For example, carlumab, a human IgG1k anti-CCL2 mAb, did not show antitumor activity as a single agent in metastatic castration-resistant prostate cancer patients in a phase II study.<sup>323</sup> Overall, the above results suggest that the CCL2-CCR2 axis may be a promising target for cancer treatment, especially when used in combination with traditional cancer-killing strategies or immunotherapies. However, the successful pharmaceutical development of CCL2/CCR2 blockers still has a long way to go.

With regard to CCL5-CCR5 signaling, systemic treatment of mice with CCL5-directed antibodies inhibit colon cancer growth, lung metastasis, and peritoneal dissemination.<sup>324</sup> CD45-immunoreactive cells in tumor tissues and adjacent healthy tissues have been found to be upregulated following CCL5 blockade. Meanwhile, CCL5 neutralization renders colon tumors more sensitive to PDGFR $\beta$ -targeted therapy.<sup>324</sup> CCL5-targeted blockade has also been demonstrated to be effective in several malignancies including breast,<sup>325</sup> gastric,<sup>326</sup> and prostate cancers.<sup>327</sup> Concurrently, much attention has been paid to the discovery of CCR5 antagonists. Maraviroc, an FDA-approved CCR5 antagonist, has been applied as an antiretroviral therapy strategy for HIV infection.<sup>328</sup> Recent studies have demonstrated its efficacy to suppress cancer cell invasiveness in a variety of cancers. Maraviroc administration resulted in cell-cycle arrest and apoptosis in both colorectal and breast cancer cell lines in vitro.325,329 In gastric cancer, treatment with maraviroc effectively reduced tumor burden, inhibited peritoneal dissemination, and prolonged the survival period.<sup>326</sup> It was also observed that maraviroc significantly inhibited pancreatic cancer liver metastasis, accompanied by marked cell-cycle arrest at the G1/S checkpoint.<sup>330</sup> TAK779 is another synthetic CCR5 antagonist that was initially developed for HIV treatments.<sup>331</sup> Menu et al. demonstrated that TAK779 has direct inhibitory effects on melanoma growth in vitro and in vivo.<sup>332</sup> Another study suggested that TAK779 could efficiently block CCR-5-dependent T-cell migration.<sup>333</sup> In a pancreatic cancer mouse model, TAK779 administration resulted in significant suppression of Treg recruitment and cancer growth.<sup>334</sup> Compared to synthetic CCR5 inhibitors, aibamine is the first natural CCR5 antagonist with an  $IC_{50}$ of 1  $\mu$ M.<sup>335</sup> Alibamine was found to suppress the invasion and metastasis of prostate cancer cells in mice.<sup>327</sup> By analyzing the chemical structure differences between anibamine and other CCR5 inhibitors, the main difference has been shown to consist of the side chains of anibamine being simple, undecorated, and aliphatic chains.<sup>336</sup> Therefore, further drug development based on the structure of anibamine may bring a novel candidate drug for cancer treatment.

CXCL12-CXCR4 is the most commonly overexpressed signaling pathway in a variety of cancers and a number of small-molecule drugs and peptide inhibitors that target CXCR4 have been developed. AMD3100 is the first CXCR4 antagonist that is used for hematopoietic stem cell mobilization during transplantation in patients with non-Hodgkin's lymphoma or multiple myeloma.<sup>337</sup> Furthermore, AMD3100 and its derivate, AMD3465, are also capable of mobilizing cancer cells in the bone marrow to ultimately enhance the efficacies of conventional therapies.<sup>338</sup> AMD3100 treatment in leukemic mice induced a ninefold increase in circulating leukemic cells.<sup>339</sup> The combined treatment of chemotherapy and AMD3100 led to a reduced tumor burden and improved OS compared with those in mice received chemotherapy alone. A similar phenomenon was also observed in a phase 1/2 study of refractory acute myeloid leukemia.<sup>339</sup> In addition to its chemosensitizing effects, AMD3100 synergistically interacts with antibodies targeting PD-L1 and CTLA-4 results in enhanced T-cell infiltration into

Chemokine-chemokine receptor targeting agents in clinical trials	
TABLE 2	

alTrials.gov First fier received		0992186 2009-10-09	1015560 2009-11-08	3473925 2018-03-22	1413022 <sup>319</sup> 2011-08-09	2345408 2015-01-26	3177187 2017-06-06	2001974 <sup>354</sup> 2013-12-05	3161431 2017-05-19		2013-02-07	3274804 2017-09-07	1736813 2012-11-29	3838367 2019-02-12	(Continue
Clinic Identi		NCT0(	NCT01	NCT0	NCT01	NCT02	NCT0	NCT02	NCT0		NCT01	NCT0	NCT01	NCT0	
Treatment		Monotherapy	Monotherapy	With pembrolizumab	With FOLFIRINOX chemotherapy	With FOLFIRINOX chemotherapy	With enzalutamide	With paclitaxel	With pembrolizumab		Monotherapy	With Pembrolizumab	Monotherapy	With carboplatin	
Tumor type		Prostate cancer	Metastatic cancer	Advanced/Metastatic solid tumors	Locally advanced or borderline resectable pancreatic cancer	Pancreatic cancer	Metastatic castration-resistant prostate cancer	Metastatic breast cancer	Metastatic melanoma		Hematologic malignancy	Metastatic colorectal cancer	Colorectal cancer patients with liver metastases	Triple negative breast neoplasms	
Phase/ status		2 <sup>a</sup>	2 <sup>a</sup>	Sb	lb <sup>a</sup>	$1b^b$	$1/2^{b}$	<b>1</b> <sup>a</sup>	$1^{\mathrm{b}}$		2 <sup>a</sup>	<b>1</b> <sup>a</sup>	<b>1</b> <sup>a</sup>	1b/2 <sup>b</sup>	
Sponsors		Centocor Research & Development	Southwest Oncology Group	Merck Sharp & Dohme	Washington University School of Medicine and National Cancer Institute	ChemoCentryx	Institute of Cancer Research, UK	Dompé Farmaceutici S.p.A	Syntrix Biosystems		Abramson Cancer Center of the University of Pennsylvania	University Hospital Heidelberg	National Center for Tumor Diseases, Heidelberg	CytoDyn	
Drug/description	CCL2-CCR2 inhibitor	Carlumab/CNTO 888 (anti-CCL2 antibody)	MLN1202/S0916 (anti-CCR2 antibody)	Navarixin/MK-7123/ SCH 527123(CXCR1 and CCR2 antagonist)	PF-04136309/PF6309 (CCR2 antagonist)	CCX872-B (CCR2 antagonist)	AZD5069 (CCR2 antagonist)	Reparixin (CXCR1 and CCR2 antagonist)	SX-682(CXCR1 and CCR2 antagonist)	CCL5-CCR5 inhibitor	Maraviroc/Celsentri (CCR5 antagonist)			Leronlimab/PRO-140 (anti-CCR5 antibody)	

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Drug/description	Sponsors	Phase/ status	Tumor type	Treatment	ClinicalTrials.gov Identifier	First received
CXCL12-CXCR4 inhibitor						
Mogamulizumab/ KW-0761 (anti-CCR4 antibody)	Kyowa Kirin	3b	Cutaneous T-cell lymphoma	Monotherapy	NCT01728805 <sup>355</sup>	2012-11-20
	European Organisation for Research and Treatment of Cancer	2 <sup>b</sup>	Stage IB-IIB cutaneous T-cell lymphoma	With radiation (total skin electron beam therapy)	NCT04128072	2019-10-16
	Kyowa Kirin	2 <sup>a</sup>	Adult T-cell leukemia-lymphoma	Monotherapy	NCT00920790 <sup>356</sup>	2009-06-15
	Kyowa Kirin	2 <sup>a</sup>	Peripheral T/NK-cell lymphoma	Monotherapy	NCT01192984	2010-09-01
	Kyowa Kirin	$1/2^{a}$	Advanced solid tumors	With nivolumab	NCT02705105	2016-03-10
	National Cancer Institute	1/2 <sup>b</sup>	Diffuse large B cell lymphoma	With pembrolizumab	NCT03309878	2017-10-16
	Kyowa Kirin	<b>1</b> <sup>a</sup>	Non-small cell lung cancer	With docetaxel	NCT02358473	2015-02-09
	Kyowa Kirin	1ª	Adult T-cell leukemia-lymphoma and peripheral T-cell lymphoma	Monotherapy	NCT00355472 <sup>357</sup>	2006-07-24
Ulocuplumab/BMS-936564/MDX- 1338 (Anti-CCR4 antibody)	Bristol-Myers Squibb	13	Acute myelogenous leukemia and selected B-cell cancers	Monotherapy	NCT01120457	2010-05-11
	Bristol-Myers Squibb	1ª	Multiple myeloma	With Revlimid/ Velcade and dexamethasone	NCT01359657	2011-05-25
AMD3100/Plerixafor (CXCR4 antagonist)	Massachusetts General Hospital	2 <sup>b</sup>	Head and neck cancer	With pembrolizumab	NCT04058145	2019-08-15
	Washington University School of Medicine	2 <sup>a</sup>	Hematologic neoplasms	With leukopheresis	NCT00914849 <sup>358</sup>	2009-06-05
	Washington University School of Medicine	1/2 <sup>a</sup>	Acute myeloid leukemia	With mitoxantrone, etoposide, and cytarabine	NCT00512252 <sup>359</sup>	2007-08-07
LY2510924 (CXCR4 antagonist)	M.D. Anderson Cancer Center	1b <sup>a</sup>	Relapsed or refractory acute myeloid leukemia	With idarubicin and cytarabine	NCT02652871	2016-01-12
	Eli Lilly and Company	2 <sup>a</sup>	Extensive stage small cell lung carcinoma	With carboplatin and etoposide	NCT01439568 <sup>360</sup>	2011-09-23
						(Continues)

TABLE 2 (Continued)

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		Phase/			<b>ClinicalTrials.gov</b>	First
Drug/description	Sponsors	status	Tumor type	Treatment	Identifier	received
Motixafortide/BKT140/BL-8040 (CXCR4 antagonist)	Biokine Therapeutics	1/2a <sup>a</sup>	Multiple myeloma	Monotherapy	NCT01010880	2009-11-10
	M.D. Anderson Cancer Center	2b <sup>b</sup>	Metastatic and recurrent pancreatic adenocarcinoma	With pembrolizumab	NCT02907099	2016-09-20
PTX-9908 (CXCR4 antagonist)	TCM Biotech International Corp.	1/2 <sup>b</sup>	Hepatocellular carcinoma	Monotherapy	NCT03812874	2019-01-23
USL-311 (CXCR4 antagonist)	Proximagen	1/2 <sup>b</sup>	Solid tumors and relapsed/recurrent GBM	Monotherapy or with lomustine	NCT02765165	2016-05-06
Balixafortide/POL6326 (CXCR4 antagonist)	Polyphor	3 <sup>b</sup>	HER2- locally recurrent or metastatic breast cancer	With Eribulin	NCT03786094	2018-12-24
	Polyphor	2 <sup>a</sup>	Multiple myeloma	Monotherapy	NCT01105403	2010-04-16
	Polyphor	$1^{a}$	Metastatic breast cancer	With Eribulin	NCT01837095 <sup>361</sup>	2013-04-22
GMI-1359 (CXCR4 antagonist)	GlycoMimetics	1b <sup>b</sup>	HR <sup>+</sup> metastatic breast cancer	Monotherapy	NCT04197999	2019-12-13
TG-0054 (CXCR4 antagonist)	TaiGen Biotechnology	2 <sup>a</sup>	Multiple myeloma and non-Hodgkin lymphoma and Hodgkin disease	With G-CSF	NCT02104427	2014-04-04
<i>ource</i> : clinicaltrials.gov.						

<sup>a</sup>Completed stage. <sup>b</sup>Ongoing stage, including both recruiting and nonrecruiting stages. tumor tissues, yielding a greater anticancer response.<sup>340</sup> Another CXCR4 antagonist, LY2510924, is also able of suppressing cancer growth and metastasis in multiple preclinical models.<sup>341,342</sup> Notably, LY2510924 was demonstrated to be clinically safe and well-tolerated in advanced solid cancers including colorectal, lung, breast, and prostate cancers.<sup>343</sup> Meanwhile, several CXCR4 smallmolecule antagonists-such as BKT140, PRX177561, and POL5551-are also capable of inhibiting cancer growth and metastasis.<sup>344–346</sup> Recent studies have also suggested that peptide inhibitors targeting the amino-terminal region of CXCR4 are also effective in inhibiting cancer growth, such as T22, TN14003, and CTCE-9908.<sup>347–349</sup> In preclinical models, TN14003 significantly inhibited pulmonary metastasis of melanomas, as well as breast cancers.<sup>350,351</sup> Similarly, CTCE-9908 showed considerable inhibitory effects on cancer growth and metastasis in malignancies including osteosarcomas and melanomas, as well as breast and prostate cancers.<sup>347,352,353</sup> These results suggest that disrupting the CXCL12-CXCR4 axis represents a viable strategy for further anticancer drug developments. However, most of the above-mentioned studies were carried out in the early course of cancer progression, and preclinical models may not fully simulate the clinical pathological process. Therefore, further clinical trials investigating the efficacies and safeties of CXCR4 antagonists are urgently needed in the context of advanced cancers.

#### 5.2 | Agents targeting tyrosine kinases

Given the activated CSF-1/CSF-1R signaling in multiple malignancies and their significant implications in predicting poor prognoses, developing agents to block the CSF-1/CSF-1R axis has attracted increasing research attention. Currently, both CSF-1/CSF-1R antibodies and CSF-1R kinase inhibitors have been successfully developed for therapeutic regulation of macrophages. The key advantage of antibodies over kinase inhibitors is their higher selectivity and safety. Currently, several CSF-1/CSF-1R targeted antibodies have been developed. H27K15 is a novel competitive monoclonal antibody for CSF-1 blockage that has been generated for cancer immunotherapy. Enzymelinked immunosorbent assays (ELISAs) have shown that H27K15 produces no cross-reaction with other tyrosine kinases and can inhibit CSF-1R phosphorylation by blocking CSF-1.<sup>362</sup> In preclinical models, H27K15 can decrease the secretion of both MCP-1 and IL-6, which play important roles in recruiting M2-like macrophages. H27K15 is also capable of preventing monocyte differentiation into CD163<sup>+</sup>CD64<sup>+</sup> M2-like macrophages and induces their differentiation into CD14<sup>-</sup>CD1a<sup>+</sup> dendritic cells.<sup>363</sup> RG7155 is designed as a humanized monoclonal antibody against

CSF-1R, with an IC<sub>50</sub> of 0.3 nM.<sup>364</sup> RG7155 can suppress CSF-1R dimerization and therefore interrupt the formation of the CSF1/CSF1-R complex. Consequently, RG7155 results in a significant reduction of CD68/CD163 macrophages.<sup>363</sup> It has been reported that objective responses were observed in 83% of patients with diffusetype tenosynovial giant-cell tumors (dt-GCTs) following RG7155 treatment, even during a short-course treatment in patients who are unable to undergo surgery.<sup>365</sup> Additionally, RG7155 is also effective in increasing the CD8/CD4 ratio in multiple solid malignancies.<sup>363,364</sup> FPA008 is also a monoclonal antibody that disrupts the binding of CSF-1R to its ligands and thus suppressing downstream receptor activation. The modulatory effects of FPA008 on macrophages and other immune cells in a variety of cancers are currently under evaluation by phase-Ia/Ib clinical trials.<sup>363</sup> With regard to adverse effects, increases in shortlived enzymes (e.g., AST, LDH) are detected, which might be due to spontaneous inhibitory effects of antibodies on CSF-1R<sup>+</sup> Kupffer cells.<sup>366</sup> Furthermore, common adverse effects also include fatigue, asthenia, facial edema, rash, and pruritus, but most of the events are only of grade 1 or 2 severities.

In addition to antibody therapies, multiple CSF-1/CSF-1R inhibitors are under development at clinical stages (Table 3). JNJ-40346527 is a selective inhibitor of CSF-1R with an IC<sub>50</sub> of 3.2 nM. In a phase-I/II clinical study of relapsed or refractory Hodgkin lymphoma, one patient out of 21 showed complete remission after JNJ-40346527 treatment (150 mg/day) for approximately 1 year. Furthermore, 11 patients remained with disease stabilization.<sup>367</sup> Notably, in vivo pharmacological activity showed that CSF-1R phosphorylation was inhibited by 95% following JNJ-40346527 treatment (150 mg/day). Meanwhile, higher doses (150-600 mg/day) also demonstrated acceptable tolerability and safety in healthy volunteers.<sup>363</sup> GW2580 is an orally bioavailable CSF-1R inhibitor with an  $IC_{50}$  of 60 nM.368 However, GW2580 shows limited inhibitory effects on macrophage recruitment and tumor growth. Priceman et al.369 treated mice to test the effects of GW2580 in blocking TAMs infiltration into growing tumors. Despite the concentration reaching as high as 160 mg/kg, only a slight decrease in TAMs was observed, while no notable changes in monocytes infiltration levels were observed in peripheral tissues including bone marrow, blood, and the spleen. Nevertheless, the combination treatment of GW2580 and VEGFR2 antibody efficiently abrogated the recruitment of TAMs and monocytic MDSCs and synergistically enhanced the antiangiogenesis response.369 Meanwhile, GW2580 was also found to increase gemcitabine therapeutic efficacy in pancreatic cancer compared to that of GW2580 alone.370 GW2580 was found to inhibit gemcitabine-induced CSF-1 overproduction in cancer cells

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Drug/description	Sponsors	status	Tumor type	Treatment	Identifier	received
CSF-1R/CSF-1 inhibitor						
Emactuzumab/RG7155/1 (anti- CSF1R antibody)	M.D. Anderson Cancer Center	2 <sup>b</sup>	Ovarian, fallopian tube, or primary peritoneal cancer	With paclitaxel and bevacizumab	NCT02923739	2016-10-05
	Hoffmann-La Roche	$1^{\mathrm{b}}$	Advanced or metastatic solid cancers	With atezolizumab	NCT02323191	2014-12-23
	Hoffmann-La Roche	1 <sup>a</sup>	Advanced solid tumors	With RO7009789	NCT02760797	2016-05-04
	Hoffmann-La Roche	1 <sup>a</sup>	Advanced solid tumors	Monotherapy	NCT01494688 <sup>378</sup>	2011-12-19
FPA008/Cabiralizumab (anti-CSF1R antibody)	Bristol-Myers Squibb	2 <sup>b</sup>	Advanced pancreatic cancer	With nivolumab	NCT03336216	2017-12-08
	Five Prime Therapeutics	1/2 <sup>b</sup>	Tenosynovial giant cell tumor	Monotherapy	NCT02471716	2015-06-15
	Five Prime Therapeutics	1a/b <sup>b</sup>	Advanced lung cancer, head and neck cancer, pancreatic cancer, ovarian cancer, renal cell carcinoma, and malignant glioma	With nivolumab	NCT02526017	2015-08-18
	Bristol-Myers Squibb	$1^{a}$	Advanced malignancies	With nivolumab	NCT03158272	2017-05-18
LY3022855/IMC-CS4 (anti-CSF1R antibody)	Eli Lilly and Company	la/b <sup>a</sup>	Solid tumor	With durvalumab or tremelimumab	NCT02718911	2016-03-24
	Dana-Farber Cancer Institute	1/2 <sup>b</sup>	Melanoma	With vemurafenib and cobimetinib	NCT03101254	2017-04-05
	Sidney Kimmel Comprehensive Cancer Center at Johns Hopkins	1 <sup>b</sup>	Pancreatic cancer	With GVAX, cyclophosphamide, and pembrolizumab	NCT03153410	2017-05-15
	Eli Lilly and Company	$1^{a}$	Advanced breast or prostate cancer	Monotherapy	NCT02265536	2014-10-16
	Eli Lilly and Company	$1^{a}$	Advanced solid tumors	Monotherapy	NCT01346358	2011-05-03
AMG820 (anti-CSF1R antibody)	Amgen	$1b/2^{a}$	Pancreatic cancer, colorectal cancer and non-small cell lung cancer	With pembrolizumab	NCT02713529	2016-03-18
	Amgen	$1^{a}$	Advanced solid tumors	Monotherapy	NCT01444404	2011-09-30
						(Continues)

TABLE 3 (Continued)						
		Phase/			Clinical trials.gov	First
Drug/description	Sponsors	status	Tumor type	Treatment	Identifier	received
MCS110 (anti- CSF1 antibody)	Yonsei University	$2^{\mathrm{b}}$	Squamous cell carcinoma	With PDR001	NCT03785496	2018-12-24
	Seoul National University Hospital	2 <sup>b</sup>	Gastric cancer	With PDR001	NCT03694977	2018-10-03
	Novartis Pharmaceuticals	2 <sup>b</sup>	Advanced triple-negative breast cancer	With carboplatin and gemcitabine	NCT02435680	2015-05-06
	Dana-Farber Cancer Institute	1/2 <sup>b</sup>	Melanoma	With dabrafenib and trametinib	NCT03455764	2018-03-07
	Novartis Pharmaceuticals	1b/2 <sup>b</sup>	Triple-negative breast cancer, pancreatic carcinoma, melanoma, and endometrial carcinoma	With PDR001	NCT02807844	2016-06-21
JNJ-40346527 (CSF-1R kinase inhibitor)	OHSU Knight Cancer Institute	2 <sup>b</sup>	Relapsed/Refractory acute myeloid leukemia	Monotherapy	NCT03557970	2018-06-15
	Janssen Research & Development	$1/2^{a}$	Relapsed or refractory Hodgkin lymphoma	Monotherapy	NCT01572519	2012-04-06
	M.D. Anderson Cancer Center	1 <sup>b</sup>	Prostate cancer	Monotherapy	NCT03177460	2017-06-06
Pexidartinib/PLX3397 (CSF-1R kinase inhibitor)	Daiichi Sankyo	3 <sup>b</sup>	Pigmented villonodular synovitis or giant cell tumor of the tendon sheath	Monotherapy	NCT02371369 <sup>379</sup>	2015-02-25
	Daiichi Sankyo	2 <sup>a</sup>	Hodgkin lymphoma	Monotherapy	NCT01217229	2010-10-08
	Daiichi Sankyo	1/2 <sup>b</sup>	Melanoma	Monotherapy	NCT02975700	2016-12-29
	National Cancer Institute	1/2 <sup>b</sup>	Plexiform neurofibroma, precursor cell lymphoblastic leukemia lymphoma and acute prolymphocytic leukemia	Monotherapy	NCT02390752	2015-03-18
	Hope Rugo, MD	$1b/2^{a}$	Metastatic breast cancer	With Eribulin	NCT01596751	2012-05-11
	Daiichi Sankyo	1b/2 <sup>b</sup>	Glioblastoma	With radiation therapy and temozolomide	NCT01790503	2013-02-13
	Daiichi Sankyo	$1/2^{a}$	Acute myeloid leukemia	Monotherapy	NCT01349049 <sup>380</sup>	2011-05-06
	Gulam Manji	1 <sup>b</sup>	Sarcoma and malignant peripheral nerve sheath tumors	With sirolimus	NCT02584647	2015-10-22
	Plexxikon	$1^{a}$	Advanced solid tumors	With Paclitaxel	NCT01525602	2012-02-03
	Daiichi Sankyo	1 <sup>b</sup>	Solid tumor	Monotherapy	NCT01004861 <sup>373</sup>	2009-10-30
	Centre Leon Berard	1 <sup>a</sup>	Advanced and metastatic colorectal cancer and pancreatic cancer	With durvalumab	NCT02777710	2016-05-19
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		Phase/			<b>Clinical trials.gov</b>	First
Drug/description	Sponsors	status	Tumor type	Treatment	Identifier	received
ARRY-382 (CSF-1R kinase inhibitor)	Array Biopharma	1ª	Metastatic cancer	Monotherapy	NCT01316822	2011-03-16
	Array BioPharma	$1b/2^{a}$	Advanced solid tumors	With pembrolizumab	NCT02880371	2016-08-26
BLZ945 (CSF-1R kinase inhibitor)	Novartis Pharmaceuticals	1/2 <sup>b</sup>	Advanced solid tumors	With PDR001	NCT02829723	2016-07-12
DCC-3014 (CSF-1R kinase inhibitor)	Deciphera Pharmaceuticals	1/2 <sup>b</sup>	Malignant solid tumors and tenosynovial giant cell tumor	Monotherapy	NCT03069469	2017-03-03
ENMD-2076 (CSF-1R kinase inhibitor)	University Health Network, Toronto	2# <sup>a</sup>	Ovarian clear cell carcinoma	Monotherapy	NCT01914510	2013-08-02
	University Health Network, Toronto	2 <sup>a</sup>	Advanced/metastatic soft tissue sarcoma	Monotherapy	NCT01719744	2012-11-01
	CASI Pharmaceuticals	$2^{\mathrm{a}}$	Advanced adult hepatocellular carcinoma and advanced fibrolamellar carcinoma	Monotherapy	NCT02234986	2014-09-09
	CASI Pharmaceuticals	2ª	Triple-negative breast cancer	Monotherapy	NCT01639248 <sup>381</sup>	2012-07-12
RON inhibitor						
Narnatumab/ IMC-RON8 (anti-RON antibody)	Eli Lilly and Company	1ª	Solid tumors	Monotherapy	NCT01119456	2010-05-07
ASLAN002/BMS 777607 (RON kinase inhibitor)	Bristol-Myers Squibb	$1/2^{a}$	Advanced or metastatic solid tumor	Monotherapy	NCT00605618	2008-01-31
	Aslan Pharmaceuticals	$1^{a}$	Advanced or metastatic solid tumor	Monotherapy	NCT01721148 <sup>382</sup>	2012-12-04
Crizotinib/ PF-02341066 (RON kinase inhibitor)	UNICANCER	2 <sup>b</sup>	Hematologic cancers and solid tumors and metastatic cancer	Monotherapy	NCT02034981 <sup>383</sup>	2014-01-14
	Dana-Farber Cancer Institute	$1^{\mathrm{b}}$	Castration-resistant prostate cancer	With enzalutamide	NCT02207504	2014-08-04
Foretinib/GSK1363089 (RON kinase inhibitor)	NCIC Clinical Trials Group	2 <sup>a</sup>	Recurrent breast cancer	Monotherapy	NCT01147484 <sup>384</sup>	2010-06-22
	GlaxoSmithKline	2 <sup>a</sup>	Head and neck cancer	Monotherapy	NCT00725764	2008-07-30
	GlaxoSmithKline	2 <sup>a</sup>	Papillary Renal-Cell Carcinoma	Monotherapy	NCT00726323 <sup>385</sup>	2008-07-31
	GlaxoSmithKline	2 <sup>a</sup>	Metastatic gastric carcinoma	Monotherapy	NCT00725712 <sup>386</sup>	2008-07-30
	GlaxoSmithKline	$1^{a}$	Solid Tumors	Monotherapy	NCT00742131	2008-08-27
	GlaxoSmithKline	1 <sup>a</sup>	Hepatocellular carcinoma	Monotherapy	NCT00920192 <sup>387</sup>	2009-06-15
<i>Source</i> : clinicaltrials.gov. <sup>a</sup> Completed stage. <sup>b</sup> Ongoing stage, including both 1	ecruiting and nonrecruiting stages.					

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and thus inhibited the recruitment of immunosuppressive myeloid cells into tumors.<sup>370</sup>

Apart from the above highly selective CSF-1R inhibitors, several CSF-1R inhibitors with lower specificity have been identified, which also exhibit broader inhibitory activity on other kinases (Table 3). PLX3397 represents one such inhibitor and has an IC<sub>50</sub> of 13 nM on CSF-1R. Additionally, PLX3397 also exhibits activities against KIT, FLT3, and PDGFR<sub>\$6,364,371,372</sub> PLX3397 displays anticancer effects in multiple malignancies. In patients with tenosynovial giant-cell tumors, 52% of cases significantly responded to PLX3397 treatment with the disease control period lasting more than 8 months.<sup>373</sup> In glioblastoma, PLX3397 not only prolonged the survival time but also slowed the disease progression rate.<sup>374</sup> PLX3397 can easily cross the bloodbrain barrier and reach glioblastoma tissue, resulting in a significant reduction of macrophage numbers and invasiveness of cancer cells. In a phase-II study of recurrent glioblastoma, PLX3397 treatment at a daily dose of 1000 mg showed acceptable tolerability and led to both CSF-1R activity inhibition and TAMs infiltration reduction in tumor tissues.<sup>372</sup> At present, multiple clinical trials are being designed and conducted to test the clinical efficacies of CSF-1R inhibitors on cancer inhibition. However, CSF-1R blockade alone usually achieves marginal therapeutic benefit, leading to the delay of tumor growth at most. Therefore, more clinical trials are of an urgent need to evaluate the combined effects of CSF-1/CSF-1R blockage and other treatment approaches including immunotherapies and standard treatment modalities. Particularly, results for combinations of CSF-1/CSF-1R blockage with immunecheckpoint inhibitors or other immunotherapeutic strategies are eagerly expected.

Several investigators have focused on the development of inhibitors targeting RON receptors (Table 3). The smallmolecule ASLAN002 is a selective inhibitor of RON receptor with an IC<sub>50</sub> < 500 nM.<sup>375</sup> ASLAN002 can block MSP-induced phosphorylation of RON. In a PyMT-MSP breast cancer mouse model, ASLAN002 administration (50 mg/kg) remarkably inhibited lung metastasis by twoto threefold.<sup>376</sup> Even when ASLAN002 was administrated after the formation of metastatic colonization, a four-fold inhibitory response was observed, indicating that RON inhibition might be a novel therapeutic option to inhibit metastatic growth as an adjuvant treatment. A mechanistic study found that ASLAN002 resulted in a proinflammatory milieu in the TME, particularly in terms of an increase in TNF- $\alpha$ -positive macrophages.<sup>376</sup> Additionally, CD8<sup>+</sup> T cells act as critical factors influencing the efficacy of ASLAN002. The anti-tumor colonization ability of ASLAN002 was greatly suppressed in the absence of cytotoxic CD8<sup>+</sup> T cells, indicating the immuneregulating cells may be primarily responsible for the anticancer effects of ASLAN002. However, future research is necessary to explore the synergistic effects between ASLAN002 and immune-checkpoint inhibitors. Another study revealed that in breast tumors harboring PIK3CA mutations, the anticancer effects of ASLAN002 were greatly limited, which were overcome by treatment with the PI3K inhibitor, NVP-BKM120.<sup>377</sup> Meanwhile, concurrent inhibition of RON and PI3K provided a more durable response in PIK3CA-wild-type breast cancer. These results suggest that further investigation focusing on the combined applications of small-molecule inhibitors is worthwhile for improving clinical prognoses of cancer patients.

#### 5.3 | Bisphosphonates

Bisphosphonates are a family of inhibitors of osteoclastmediated bone resorption and are widely applied in the treatment of bone diseases including postmenopausal osteoporosis, Paget disease, and bone metastasis. The first generation of nonamino bisphosphonates, such as etidronate and clodronate, are metabolized intracellularly into cytotoxic adenosine-triphosphate analogs that induce osteoclast cell death.<sup>388</sup> Conversely, the second and third generations of bisphosphonates-such as ibandronate, risedronate, pamidronate, and zoledronic acid-are much more potent than nonamino compounds by inhibiting farnesyl-pyrophosphate synthases, which also results in osteoclast apoptosis.<sup>389</sup> Moreover, preclinical studies demonstrate that bisphosphonates may also exhibit extraskeletal therapeutic effects in mouse breast cancer xenografts.<sup>390</sup> For example, zoledronic acid was found to be phagocyted by TAMs and thus induced TAMs apoptosis and repolarization toward M1-like phenotype.9 In breast cancer patients, radiolabeled bisphosphonate was also found in the resected tumor infiltrated with TAMs and microcalcifications. Several studies have demonstrated the proinflammatory activity of amino-bisphosphonates in vitro. Ibandronate was found to enhance LPS-stimulated IL-6 and IL-1 $\beta$ secretion from mouse Raw264.7 macrophages.<sup>391</sup> A remarkable elevation in TNFa expression was also observed following pamidronate and zoledronic-acid administrations.<sup>392</sup> In mouse xenografts, ibandronate treatment led to increased infiltration of extravasated leukocytes and rolling leukocytes, accompanied by the stimulated production of TNF $\alpha$  and IFN- $\gamma$  in peripheral blood, indicating its proinflammatory effects in vivo.<sup>393</sup> In cancer patients treated with zoledronic acid, it was also found that the circulating concentrations of IFN- $\gamma$ , TNF $\alpha$ , and IL-6 were all significantly elevated following drug administration, further confirming that amino-bisphosphonates might promote M2-like to M1-like

repolarization to trigger inflammatory responses.<sup>394</sup> By contrast, nonamino bisphosphonates have been demonstrated to inhibit cancer progression via suppressing the growth of macrophages. Clodronate inhibits iNOS expression in Raw264.7 macrophages, accompanied by a significant reduction of IL-6, IL-1 $\beta$ , TNF $\alpha$ , and NO secretions at noncytotoxic concentrations.<sup>395</sup> Therefore, nonamino bisphosphonates might suppress macrophages in a cytostatic but not cytotoxic manner. Based on the above findings, bisphosphonates have been commonly applied to deplete TAMs in preclinical models. Although bisphosphonates have yielded improved survival in various kinds of preclinical animal models, bisphosphonates treatment alone is not able to induce complete regression of tumors. In fact, the application of bisphosphonates may induce side effects on residential macrophages in normal tissues, thereby influencing antitumor innate immunity. Considering that the complete depletion of macrophages in tumors could lead to worse therapeutic outcomes in some cases, TAMs-targeted reprogramming strategies that spare the potentially beneficial macrophages may be more preferable in the future.

# 5.4 | Nanoparticle or exosomal-targeted delivery

Although pharmaceutical inhibitors and monoclonal antibodies have been developed for TAMs-targeting treatments, their therapeutic efficacies have been greatly limited by difficulties in penetrating biological barriers. Progress in nanotechnology has provided a valuable opportunity for improving drug-loading/drug-releasing parameters, biocompatibilities, and drug-circulation time. The engineering of new nanoparticles with the ability to target and kill or reeducate TAMs has stood up as a promising strategy for cancer treatment in recent years. As an important member of the mononuclear phagocytic system, macrophages play a major role in nanomaterials recognition, uptake, processing, and clearance.<sup>396</sup> Therefore, the loading of macrophages with nanomedicines may be used to profit from the high infiltration and the innate tumor homing abilities of macrophages, allowing the drug release in the bulk of the tumor. For example, Rao et al. reported that the magnetic nanoparticles can be used to promote the repolarization of the M2-like TAMs as well as the systemic circulation and tumor accumulation of the loaded drugs.<sup>397</sup> Notably, there are still some challenges for the effective targeting of TAMs with nanoparticles and their application in the clinic. The major one can be attributed to the undesirable clearance of nanoparticles by the mononuclear phagocyte system (macrophages) in clearance organs (liver, lung, or spleen) upon their intravenous injection. *In vivo* studies have demonstrated high nanomaterials sequestration by macrophages in clearance organs such as liver, spleen, and kidney.<sup>396</sup> This is because that the immune system may recognize the components of nanoparticles (e.g., shell, core, surface-decorating moieties, and cargoes) as foreign, and initiates an immune response through a complex process.<sup>398</sup> Therefore, because of the synthetic origin, nanoparticles are usually recognized by the host immune system and are preferentially sequestered by macrophages in filtering organs.

Recently, exosomes have emerged as natural nano-sized vesicles for drug delivery. Given their nanoscale size, excellent biocompatibility, potential capacity to express targeting ligands, and natural capacity to carry macromolecules, the use of exosomes for cancer treatment has raised considerable interest.<sup>399,400</sup> Due to the lipid bilayered membrane, exosomes exhibit good permeability and can cross most biological membranes, including the bloodbrain barrier.<sup>401</sup> After reaching recipient cells, exosomes can activate certain signaling pathways by fusion to the plasma membrane and subsequent secretion of the contents into the cytoplasm of recipient cells. Alternatively, exosomes can enter cells through endocytic processes including receptor-mediated endocytosis, micropinocytosis, and phagocytosis. Moreover, exosomes have significant advantages in drug delivery areas without inducing proinflammatory responses or adverse immune reactions. More importantly, the composition of exosomal surfaces permits modifications for enhancing interactions between recipient cells and exosomes. Surface modification of naturally purified exosomes with GE11, an epidermal growth factor, was demonstrated to increase the targeting potential of exosomes to EGFR<sup>+</sup> breast cancer cells both in vitro and *in vivo*.<sup>402</sup> Given these priorities, exosomes represent a promising candidate tool for the delivery of chemotherapeutics, as well as nucleic acids and phytochemicals. For example, doxorubicin treatment through the exosome delivery system induced a 10-fold increase in cancer cell death when compared with free doxorubicin.<sup>403</sup> Similarly, exosome-loaded anti-miR-214 led to a remarkable inhibition of gastric tumor growth.<sup>404</sup> Interestingly, curcumin has also been shown to be encapsulated into exosomes and to interfere with colon carcinogenesis (NCT01294072).

Given that exosomes can be preferentially sequestered by macrophages,<sup>405,406</sup> it is becoming an attractive vehicle for delivering genetic drugs or cytotoxic agents to inhibit TAMs. Exosomes containing miRNA-155 and 125b2 derived from pancreatic cancer cells induced macrophage differentiation to the M1-like phenotype.<sup>407</sup> Similarly, exosomes transfected with wt-p53 and miRNA-125b secreted from colon cancer cells also mediated macrophage repolarization toward the M1-like phenotype.<sup>408</sup> Interestingly, epigallocatechin-gallate-treated breast cancer cells-derived exosomes were also found to impair tumor growth by suppressing TAMs infiltration and M2-like phenotype polarization.<sup>137</sup> Recently, a commercialized engineered exosome carrying antisense oligonucleotide (exoASO) was developed by Codiak Bioscience. It was designed to deliver ASO to TAMs and to specifically inhibit the expression of immune-suppressive transcription factors, including STAT6 and C/EBPT.<sup>137</sup> Preclinical studies demonstrate that exoASO effectively reprograms M2-like macrophages to a pro-inflammatory M1-like phenotype, promoting targeted anti-tumor activity in vivo. In vitro, exoASO is preferentially taken up by M2-like macrophages to a significantly greater extent than that of free ASO, resulting in a greater decrease of STAT6 and C/EBPß mRNA levels. Subsequent gene expression analysis and cytokine assays showed a 40-fold increase in TNF $\alpha$ and a 29-fold decrease in IL-10 associated with exoASO treatment, consistent with repolarization from immunosuppressive M2-like macrophages to immune-stimulatory M1-like macrophages. All of these findings suggest that exosomes may represent promising drug-delivery systems for modulating the biological functions of TAMs. Notably, there are still many difficulties to overcome before the successful application of the exosomal delivery system for TAMs modulation in the future. The most prominent one may be that different exosome isolation techniques have reported poor yield and loss of functional properties.<sup>409</sup> To solve this limitation, reengineering TAMs-targeted exosomes with the synthetic liposomes as a refined biomimetic nanostructure is suggested.<sup>409</sup>

### 5.5 | Other therapeutic strategies

In recent years, multiple novel TAMs reprogramming strategies have been reported with promising future. For example, Christopher B et al. reported that R848, an agonist of toll-like receptor 7 and 8 (TRL7/8), could significantly induce the M1-like phenotype polarization of macrophages in vitro, while the R848-loaded  $\beta$ cyclodextrin nanoparticles could promote the repolarization of TAMs toward an M1-like phenotype in multiple tumor models *in vivo*.<sup>410</sup> Feng *et al.* synthesized the novel TLR agonist of acGM using glucomannan polysaccharide with acetyl modification, which could specifically activate the TLR2 signaling and thus induce TAMs repolarization into an antitumor phenotype.<sup>411</sup> Macrophages can also be targeted via CD40 agonists to boost the antitumor response. Gregory et al. reported that the CD40 agonist FGK45 could reactivate TAMs, which became tumoricidal and thus rapidly facilitated the depletion of tumor stroma and the tumor regression of pancreatic ductal adenocarcinoma.412 Blocking checkpoints on T cells have

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recently shown unprecedented, durable cures in the clinic. A similar strategy targeting macrophages is also developing by blocking CD47 from binding its receptor signal regulatory protein- $\alpha$  (SIRP $\alpha$ ) on macrophage surface.<sup>413</sup> It has been reported that interfering of CD47-SIRP $\alpha$  interaction by blocking antibodies could promote the phagocytosis of TAMs and consequently suppresses cholangiocarcinoma growth and metastasis.<sup>414</sup> Nowadays, a number of CD47-SIRP $\alpha$  axis blockers, such as Hu5F9-G4, CC-90002, SRF231, ALX148, and IBI188, are under clinical trials.<sup>413</sup>

# 6 | CONCLUSIONS, CHALLENGES, AND PERSPECTIVES

With a deeper understanding of cancer immunology, targeting TAMs provides a novel approach to improve anticancer therapies. However, since there exists complex intercellular crosstalk involving TAMs in the TME, TAMs elimination may elicit multifaceted stromal reactions in the host that are unable to predict and may vary among patients. Additionally, some patients may be refractory to TAMs-targeting therapies due to gene mutations. For example, it has been reported that certain singlenucleotide polymorphisms in CSF-1R can decrease the therapeutic efficacy of emactuzumab.<sup>137</sup> Therefore, it is important and significant to explore the synergistic effects of TAMs-targeting approaches with other immunomodulatory strategies, especially for checkpoint-blockade-based therapies. Checkpoint inhibition unleashes the brakes, such as CTLA4 and PD-1, to enhance the effective recognition and killing of tumor cells by immune cells. However, only approximately 20% of cancer patients respond to checkpoint-inhibition therapies, where mixed responses can limit treatment effectiveness and cause drug resistance and/or distant metastasis. TAMs may act as important cellular components influencing the responsiveness of checkpoint-blockade treatments. On one hand, TAMs express PD-L1 and PD-L2, which could aggravate the immunosuppressive TME and impair cytotoxic T lymphocytes against cancer cells.<sup>14,15,415</sup> On the other hand, TAMs also expressed TREM2 (triggering receptor expressed on myeloid cells 2) that remodel the tumor myeloid landscape and weaken the responsiveness of anti-PD-1 immunotherapy.<sup>416</sup> Currently, a variety of clinical trials are ongoing to examine the synergistic effects of TAMstargeting approaches with PD-L1 or CTLA-4 blockades. Preclinical studies have also demonstrated that CSF1R antagonists can boost the efficacy of checkpoint inhibition in mouse models of pancreatic, breast, cervical, and ovarian cancers. Moreover, in immunotherapy-resistant pancreatic cancer patients, CSF-1R antagonists combined with PD-1-targeting approaches result in partial responses

in some patients. All of these findings indicate the promising future and significance of TAMs-targeting strategies in cancer immunotherapies.

With an increasing number of available drugs for modulating TAMs to induce anti-tumor responses by targeting different molecular targets, the next challenge will be how to deliver these immunoregulatory drugs into TAMs effectively and selectively while minimizing the harmful off-target side effects. Although nanovesicles, such as exosomes, provide an ideal platform for drug loading and delivery and can be primarily internalized by macrophages, their biodistributions in vivo are largely affected by residential macrophages outside of tumors, such as Kupffer cells in the liver. To improve TAMs selectivity over residential macrophages, several studies have utilized the unique physical and chemical properties of the TME to optimize their targeting systems. For example, the acid-sensitive PEG 2000 was applied to modify nanoparticles to selectively eliminate TAMs as PEG 2000 can be cleaved in the acidic TME to release cytotoxic drugs.<sup>417</sup> This therapeutic strategy can minimize the offtarget effects of PEG 2000-modified nanoparticles on resident macrophages. Additionally, targeting ligands have also been applied to improve the preferential delivery to TAMs via utilizing the specific ligand-receptor interactions. CD206 represents one of the most commonly targeted receptors for macrophage delivery because of its TAMs-specific overexpression characteristic. As the native ligand of CD206, mannose can be conjugated to nanovesicles easily and result in an enhanced uptake by TAMs.<sup>417</sup> TAMs also overexpress folate receptor  $\beta$ , the ligand of which is folic acid. Therefore, folic-acid-modified nanovesicles have been developed as an important cancertargeting strategy.<sup>418</sup> Another factor influencing the antitumor efficacy of nanovesicles is the ability of drug-loaded macrophages to infiltrate into tumors. A recent study suggested that the size of nanovesicles can significantly alter the motilities of macrophages.<sup>419</sup> Although smaller nanovesicles (30-50 nm) usually have higher uptake into macrophages compared to that of larger nanocarriers (100-500 nm), smaller nanovesicles were found to significantly retard the motilities of macrophages. This study suggested that 100-nm nanovesicles provide a good balance for effective drug loading and macrophage migration. Hence, exosomes are an ideal and promising type of nanovesicle for developing TAMs-targeting strategies.

Taken together, the pharmaceutical targets of TAMs and their corresponding drugs are rapidly being explored and developed. With the development of single-cell sequencing, various "omic" analyses, and nanotechnology, our understanding of the heterogeneity of TAMs in the tumor milieu will be considerably enhanced. It is expected that future studies will further elucidate the clinical benefits resulting from combined therapies of TAMs-targeting strategies with traditional treatments or immunotherapies. Meanwhile, a more advanced drug delivery system via exosome-like nanovesicles is anticipated to help reduce the off-target/side effects of TAMs-targeting approaches.

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#### AUTHOR CONTRIBUTIONS

NW and ZYW conceived, wrote, and revised the article. SQW drew the figures and revised the article. XW generated the tables and added references. YFZ, BWY, JPZ, BP, and JLG participated in the work of reference selection and article check.

#### **COMPETING INTERESTS**

The authors declare that they have no competing interests.

### AVAILABILITY OF DATA AND

MATERIALS

Not applicable.

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